

Characterization of heterochromatic regions in two *Triturus alpestris* subspecies (Urodela: Salamandridae)

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The fluorescence analysis carried out in *Triturus alpestris alpestris* and *Triturus alpestris cyreni* subspecies has revealed differences related to the content and distribution of AT-rich, GC-rich and non-fluorescent heterochromatic bands. These results provide new evidence on the chromosome differentiation undergone in their phylogenetic splitting.

INTRODUCTION

Fluorescence differential staining of constitutive heterochromatic regions in the chromosomes is generally assumed to be the result of the existence of clusters of highly repetitive DNA. Biochemical analysis has demonstrated that the adenine-thymine (AT) base richness induces certain fluorescent bands with some fluorochromes while the guanine-cytosine (GC) rich regions show differential fluorescence with other fluorochromes (Schmid, 1980; Schweizer, 1981). Thus these techniques are particularly useful to analyze both the DNA composition and the location of heterochromatic regions in the chromosomes.

To date ten subspecies of *Triturus alpestris* have been described on morphological grounds (Thorn, 1968; Dubois and Breuil, 1983). *T.a. alpestris* occupies almost the entire range of distribution of the species. The remaining subspecies are exclusively represented by certain isolated populations. One of these subspecies, *T.a. cyreni*, was first described by Wolstertorff (1932) on the basis of its restricted distribution. Its populations are only found in restricted habitats of northern Spain thus being separated from the remaining subspecies of *T. alpestris* by a large part of France.

In a previous study we reported that *T.a. cyreni* displays certain peculiar characteristics in the distribution of constitutive heterochromatin which are not found in *T.a. alpestris* (Herrero and Arano, 1986). In this paper we report the nature of these heterochromatic differences between the two sub-

species of *T. alpestris*, on the basis of fluorescence banding.

MATERIALS AND METHODS

Ten adult specimens of *T.a. cyreni* (five ♀ and five ♂) per population were used in this study. They were collected from Lago de la Ercina, Lagos de Somiedo and Santillana del Mar (Spain). Four adult specimens of *T.a. alpestris* (two ♀ and two ♂) were collected from Bonn (FRG).

Males and females were injected with colchicine solution (0.1 per cent) 7 h before they were sacrificed. Pieces of intestine and testes were extracted in a hypotonic KCl (0.1 M) solution at room temperature, they were fixed in 3:1 ethanol:acetic acid for 24 h. Squash preparations were made in 45 per cent acetic acid and the coverslip was removed after freezing in liquid nitrogen. Then slides were dehydrated and subjected to the C-banding technique reported by Sumner (1972).

From each individual about 50 intestinal and testicular metaphases were studied.

The fluorescence analysis was carried out using the staining methodology developed by Schweizer (1976; 1980; and personal communication). We used chromomycin A₃ (CMA₃), Distamycin (DA) and 4'-6 diamino-2-phenylindole (DAPI) dyes supplied by SERVA.

Fluorescence observations were carried out using a Zeiss Photo III microscope and photographs were taken with Valca 125 ASA film.

RESULTS

The two subspecies used in this study have 24 biallelic chromosomes. The C-banding pattern of both subspecies consists of centromeric bands in almost all chromosomes. Thin pericentromeric bands in variable numbers (single, double or triple) were observed in all chromosomes. Pairs 1, 3 and 8 display distal bands which affect both arms in pairs 1 and 3 and the longest arm of pair 8. The differences between the subspecies relate to the presence in *T.a. alpestris* of interstitial bands in pairs 4 and 6 and a distal band in pair 12 that are absent from *T.a. cyreni*. On the other hand *T.a. cyreni* presents several bands not found in *T.a. alpestris*. Some are located in the centromere (pair 8), and others in interstitial (pairs 3 and 8) or telomeric (pairs 10 and 11) positions. Some coincident bands are larger in *T.a. cyreni* than in *T.a. alpestris* (fig. 1(a) and (1b)) (Schmid *et al.*, 1979; Herrero and Arano, 1986). We have not found inter-individual variations in banding patterns.

On the other hand, while males of *T.a. alpestris* possess a characteristic XY sex chromosome pair (number 4), such differentiation has not been found in *T.a. cyreni* (Herrero and Arano, 1986). Both males and females of *T.a. cyreni* present a telomeric grey C-band in the homologues of pair 4. The application of fluorochromes to both *T. alpestris* subspecies reveals new information as regards the characteristics of the C-heterochromatic regions. Three kinds of fluorescent and non-fluorescent bands have been found:

(i) AT-rich C-bands.

These regions correspond to those that are positive to DA-DAPI (Distamycin/4'-6 diamidino-2-phenylindole). Most C-bands are included in this pattern, so that the differences found between the two subspecies correspond to the absence/presence of some C-bands, particularly in the centromeric and pericentromeric regions. Good examples are the pairs 4, 6 and 8 (figs. 1 and 2).

(ii) GC-rich C-bands.

DA-CMA₃ (Distamycin/Chromomycin A₃) reveals very restricted areas of positive fluorescence. The positive fluorescence occurs in 5 autosomes pairs (1, 3, 7, 8 and 9) in *T.a. alpestris* and is located in pairs 1, 3, 8, 9 and 10 in *T.a. cyreni*. The differences between the

two subspecies involve the positive CMA₃ fluorescence of pairs 3, 7, 8 and 10. The differences between the 3, 7 and 10 pairs are not revealed by C-banding (figs. 1 and 2).

(iii) Non fluorescent C-bands.

Such regions show a similar staining intensity to the euchromatin. This labelling has been observed in the terminal C-band of pair 12 in *T.a. alpestris* and in the interstitial and telomeric C-bands of pairs 3 and 11 in *T.a. cyreni* (figs. 1 and 2).

This analysis has not revealed more heterochromatic regions than C-banding has.

DISCUSSION

Our results show the divergence between two isolated groups and allow us to observe the initiation of a possible phylogenetic splitting. Roger's distance reveals that these subspecies have been isolated since the Pleistocene age (Arano and Arntzen, 1987). During this time new bands in *T.a. cyreni* and in *T.a. alpestris* have emerged or have been lost (depending which state were ancestral). On the other hand, four of the heterochromatic shared bands have changed their base composition, two of them being GC-rich in *T.a. alpestris* and AT-rich in *T.a. cyreni*, and the other two being GC-rich in *T.a. cyreni* and AT-rich in *T.a. alpestris*.

Two models have been recently proposed to explain the location and growth of new heterochromatin regions. One of them (Schweizer and Loidl, 1986) predicts that the telomeres are C-band initiation sites from which there is a heterochromatin transfer to interstitial sites of other non-homologous chromosomes during the mitotic and meiotic processes. This model predicts that a karyotype with chromosomes of similar length will tend to have telomeric C-bands rather than interstitial bands and a karyotype with chromosomes of different sizes and arm ratios will tend to possess interstitial and telomeric bands. None of these predictions are in accordance with the C-band patterns seen in the two subspecies studied as the telomeric bands are not predominant in them.

The second model (Macgregor and Sessions, 1986) proposes that C-bands would be dispersed from the heterochromatic initiation sites in the centromeres towards the pericentric region and from there to the telomeric region. Therefore, those karyotypes with more telomeric heterochromatin would have an older phylogenetic status. The slight trend observed in *T.a. cyreni* to have more telomeric bands than *T.a. alpestris*, where most

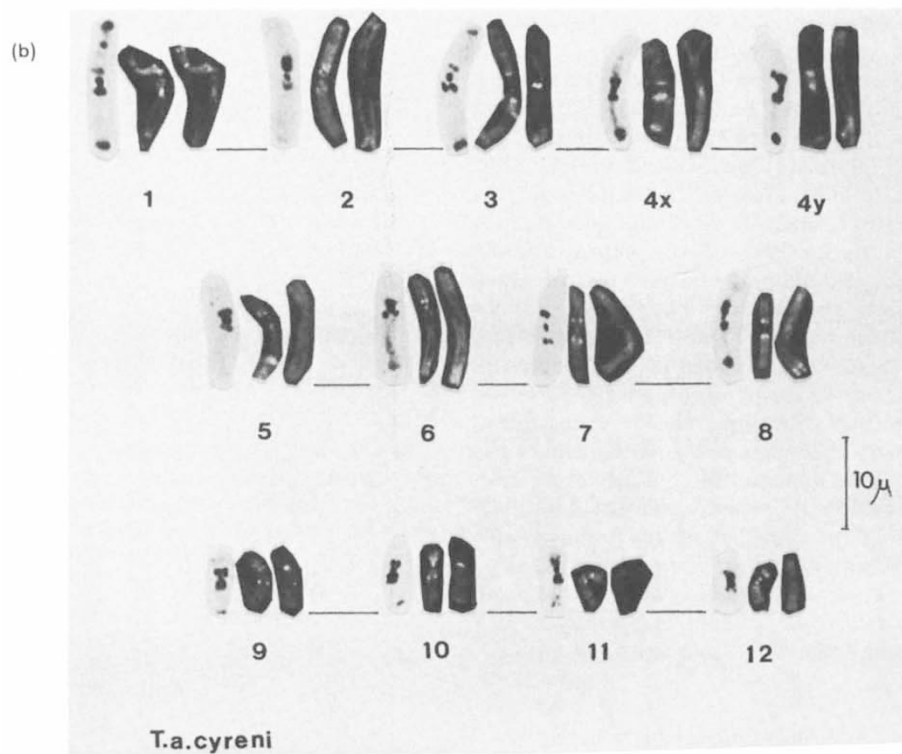
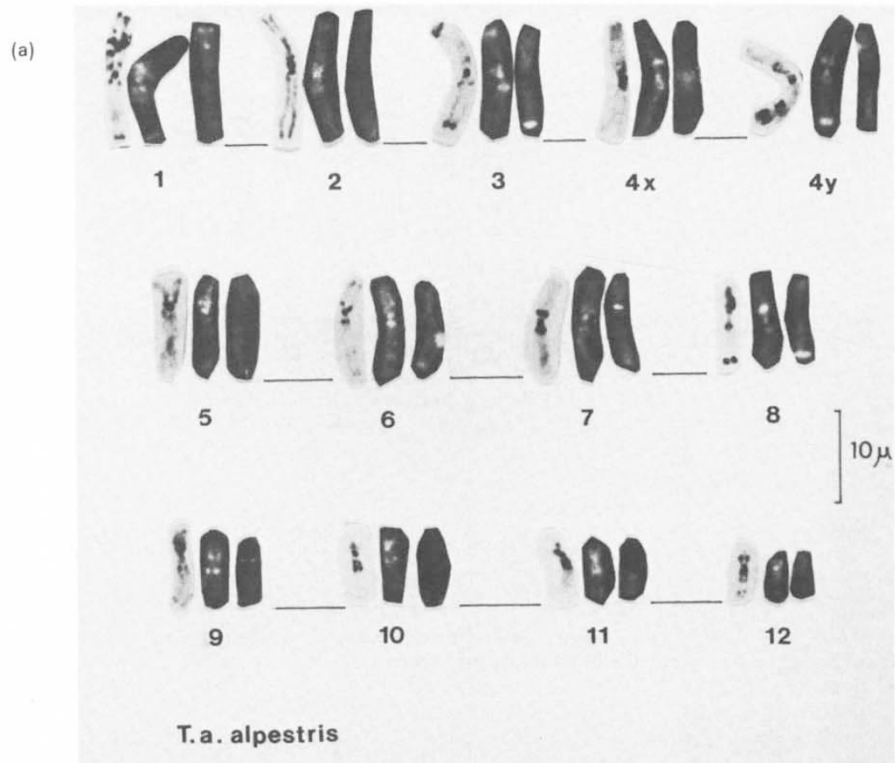


Figure 1 Haploid karyotypes of: a, *Triturus alpestris alpestris*; b, *Triturus alpestris cyreni*, after C-banding and fluorescent staining. Within each chromosomal triad, the first chromosome corresponds to the C-banding, the second one to the DA-DAPI staining and the third one to the DA-CMA₃ staining.



Figure 2 Idiograms of *Triturus alpestris alpestris* and *Triturus alpestris cyreni* showing the differential response to distinct fluorochromes of the C-bands contained in their chromosome complements.

bands are pericentric, would corroborate electrophoretic results which suggest that *T. a. cyreni* represents an older evolutionary status. However, differences in the composition of the heterochromatin of particular chromosomes might reveal a different origin for each band, which is the case for pairs 3, 9 and 10 in *T. a. cyreni*, and pairs 3, 7 and 9 in *T. a. alpestris*. Hence, this model would not fully explain the *T. alpestris* C-banding distribution and origin. It seems clear that chance could have played a major role in the causation of heterochromatic regions in the early stages of subspecies evolution and there is no evidence of a tendency for these regions to appear in predetermined locations in the chromosome complement.

Finally, our results from the comparative analysis of fluorescent banding show differences between the two subspecies not only in respect to the heterochromatic distribution, but also concerning the composition of several shared C-bands. Taking into account that C-banding patterns of *Triturus* are extremely conservative (John, 1988), the sharp differences observed between *T. a. alpestris* and *T. a. cyreni* suggest that an important divergence process may affect the *T. alpestris* complex.

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