# Localization of NORs in karyotypes of four Rana species

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#### **Abstract**

NORs were localized by Ag-AS-staining in karyotypes of Rana nigromaculata, R. macrocnemis, R. latastei (2n = 26) and R. chensinensis, 2n = 24. As in the majority of other Rana species, the NORs are located in the long arms of the 10th (or 11th in R. chensinensis) chromosome pair. The presence of a second secondary constriction in R. latastei, as reported earlier, was not confirmed.

#### Introduction

The presence and position of secondary constrictions which are potentially the nucleolus organizer regions (NORs, the sites of 18S + 28S rDNA) are important characteristics in karyosystematics. With regards to amphibian karyotypes, especially those of Rana species, secondary constrictions were discussed by Hennen (1964), Seto (1965), Guillemin (1967), Ullerich (1967), Günther (1970), Nishioka (1972), Kuramoto (1972, 1980), Kuramoto et al., (1973), Ivanov and Madyanov (1973), Belcheva and Genova (1974), Haertel et al. (1974), Popov (1974), Orlova et al. (1977), Koref-Santibanez and Günther (1980), Green et al. (1981), Green (1983). NORs can be specifically stained by the Ag-AS-method (Goodpasture & Bloom, 1975), although some other chromosome sites are stained during this procedure too (Nardi et al., 1978; Vitelli et al., 1982; Bailly, 1983).

As a rule, intensively Ag-stained regions of chromosomes correspond to secondary constrictions; in some cases, however, data published by authors who used the Ag-staining method and those who did not, were different. For instance, the results of Ag-staining of *Bufo* chromosomes (Schmid, 1978a; Beck & Mahan, 1979; Birstein,

1981a) point out that Bogart (1972) perhaps overestimated the number of secondary constrictions; in fact, NORs (and the corresponding secondary constrictions) are located in homologues of only one *Bufo* chromosome pair – the 1st, 6th or 11th. Previously chromosomes of some *Rana* species were Ag-stained (Ward, 1977; Schmid, 1978b; Belcheva et al., 1981; Birstein, 1981a, b). This paper describes the position of NORs in karyotypes of four more *Rana* species and discusses the localization of NORs in *Rana* karyotypes in general.

## Material and methods

One species of green frog, Rana nigromaculata (2\,\text{?}, 2\,\text{?} from Karamet-Niyaz, Turkmenian SSR), and three brown frog species, R. macrocnemis (1\,\text{?}, 1\,\text{?} from the Caucasian Wild Life Reserve, Guseripl), R. latastei (2\,\text{?} from the environments of Mantova, Bosco della Fontana, Italy – Orlova et al., 1977) and R. chensinensis (3\,\text{?} from the environments of Yujnosakhalinsk, Sakhalin island) were investigated. Chromosome preparations were made from bone marrow cells of colchicinized animals (Aleksandrovskaya et al., 1979). The onestage method of Ag-AS-staining was used (Howell

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& Black, 1980). One drop of gelatin solution (2% solution in water acidified with formic acid) and two drops of 50% AgNO<sub>3</sub> solution were placed on the preparation, then the drops were mixed, and the mixture was covered with a cover glass. The preparation was heated to 70 °C for 2.0-2.5 min, the cover glass taken off and the preparation washed with water, dried and analyzed. In some cases after the preparations had been photographed, silver was removed according to Warburton and Henderson (1979). The slides were first treated with 7.5% sodium ferricyanide (4 min) and after that with a 20% solution of sodium thiosulfate. They were then washed with water and stained with Giemsa.

### Results and discussion

Figure 1 shows karyotypes of the four investigated Rana species after Ag-staining of the chromosomes. The karyotypes of R. nigromaculata, R. macrocnemis and R. latastei consist of 26 chromosomes while the R. chensinensis karyotype consists of 24 chromosomes; these data coincide with findings of Nishioka (1972), Ivanov and Madyanov (1973) and Orlova et al. (1977). On the basis of relative length the chromosomes of the first three species (2n = 26) can be divided into two groups: large chromosomes (pairs 1-5) and small ones (pairs 6-13). All large chromosomes are meta- or submetacentric, while some of the small ones are subtelocentric (e.g. pairs Nos. 7, 9 in the R. latastei karyotype). The three karyotypes are very similar and the main difference between R. nigromaculata and R. macrocnemis and R. latastei is the centromeric position in the 4th pair: in R. nigromaculata these are almost subtelocentric, in R. macrocnemis and R. latastei they are submetacentric. It should be noted that we analyzed the karyotypes of R. nigromaculata from Central Asia, where this species had been introduced from China (Aleksandrovskaya & Orlova, 1981). The karyotypes of this form and of the Far Eastern R. nigromaculata are practically indistinguishable (Aleksandrovskaya & Orlova, 1981).

In all three frog species (2n = 26) NORs are located in the long arms of the homologues of the 10th chromosome pair (Fig. 1). In *R. nigromaculata* and *R. latastei* these are submetacentric and NORs are situated about the middle of the long

arms; in *R. macrocnemis* they are almost metacentric and NORs are located in the long arms near the centromeres.

Orlova et al. (1977) described two secondary constrictions (one in each arm) in the 7th pair of R. latastei. According to our numeration, this pair could be No. 10. However, we did not find any additional constriction in the small chromosomes, neither after Ag-staining nor after silver had been removed and the same preparations were stained with Giemsa. Thus, the 2nd secondary constriction described by Orlova et al. (1977), could be the result of an artefact.

There is little difference in the length of Rana chromosomes Nos 7-11 (and especially Nos. 9-11). That is why several authors gave different numbers to the chromosomes having secondary constrictions (e.g. Kuramoto, 1972, and Kuramoto et al., 1973, reported secondary constrictions in the 9th pair of the Far Eastern Rana species, instead of in the 10th). Some authors did not find constrictions at all. For instance, secondary constrictions previously were not found in R. macrocnemis (Ivanov & Madyanov, 1973), while we localized the NORs in the 10th pair of this species.

The difference in length of large and small chromosomes of R. chensinensis, 2n = 24, is not as evident as in Rana species with 2n = 26. The chromosomes grade smoothly as in the case of R. dybowskii and R. ornativentris (Green, 1983). NORs of R. chensinensis are located in the long arms of the homologues of one of the smallest pairs, which we designate by No. 10, as in another Agstained Rana species, 2n = 24, R. macrodon (Schmid, 1980a). In the related species, R. dybowskii, this pair was designated as No. 11 (Green, 1983).

The karyotype of *R. chensinensis* is very similar to that of *R. dybowskii* (Green, 1983), there are two pairs of submetacentrics, Nos. 8 and 10 (11 in Fig. 1) in these two karyotypes, which probably originated after an initial chromosome number reduction from a 26-chromosome ancestor (Green, 1983). The C-band pattern of *R. dybowskii* suggested to Green (1983) that the 24-chromosome karyotype could arise by a fusion of two small chromosome pairs in the karyotype of a 26-chromosome ancestor. A new chromosome of intermediate size (No. 6) derived from this fusion would shift the small NOR-bearing chromosome from its

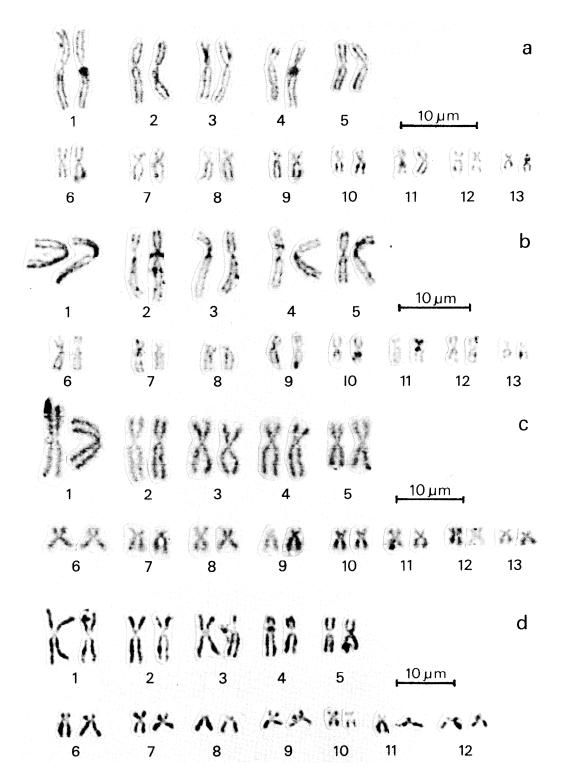


Fig. 1. Karyotypes of Rana nigromaculata (a), R. macrocnemis (b), R. latastei (c), and R. chensinensis (d). Chromosomes were stained by the Ag-AS method; the homologues of the 10th pairs have Ag-positive NORs in the long arms.

Table 1. The localization of nucleolus organizer regions in karyotypes of Ranidae (Ag-positive sites).

Species	Origin	2n	Position of Ag-NOR <sup>a</sup>	References
Rana esculenta	Central Europe	26	10q <sup>b</sup>	Schmid, 1978b; Birstein, 1981a
R. graeca	Western Europe	26	10q <sup>c</sup>	Vitelli et al., 1982
R. latastei	Western Europe	26	10q	This paper
R. lessonae	Central Europe	26	10q	Birstein, 1981a
R. ridibunda	Central Europe	26	10q	Schmid, 1978b; Belcheva et al., 1981;
				Birstein, 1981a, b
R. temporaria	Central Europe	26	10q <sup>b</sup>	Schmid, 1978b; Birstein, 1981a
R. macrocnemis	The Caucasus	26	10q	This paper
R. erythrea	Southeast Asia	26	10q (6q, 8q,	Schmid, 1978b; 1980a
			8p, 11p, 13p	
R. nigromaculata	Central Asia	26	10q	This paper
R. chensinensis	Far East	24	10q <sup>d</sup>	This paper
R. macrodon	Southeast Asia	24	10g	Schmid, 1980a
R. blairi	North America	26	10q	Ward, 1977
R. catesbeiana	North America	26	10q <sup>b</sup> (8p, 12p <sup>e</sup> )	Schmid, 1978b
R. magnaocularis	North America		10q	Ward, 1977
R. palustris	North America	26	10q (6q, 6p, 7p,	Schmid, 1978b; 1980a
			7q, 12p, 13q)	
R. sphenocephala	North America	26	10q (9p, 13p)	Schmid, 1978b
Pyxicephalus adspersus	South Africa	26	6p (8p)	Schmid, 1978a, b
P. delalandii	South Africa	26	13p	Schmid, 1980b

<sup>&</sup>lt;sup>a</sup> q: long arm; p: short arm; the additional Ag-positive sites (not NORs) are listed in brackets.

usual tenth position in the 26-chromosome karyotypes to the eleventh position in the 24-chromosome *R. chensinensis* and *R. dybowskii* karyotypes. The same shift has probably occurred in the karyotype of *R. ornativentris* (Green, 1983).

Thus NORs are located in the 10th (or in the homologous 11th) chromosome pairs of four investigated Rana species. The same is true for many other Rana species (Table 1). Only in African species of Pyxicephalus, which are closely related to Rana, are NORs located in the 6th or 13th pairs (Schmid, 1980a, b). In the case of P.adspersus, there is one additional NOR in the short arm of a small homologue of the heteromorphic pair No. 8 (Schmid, 1980b). It seems possible that this change in the P. adspersus karyotype, in comparison with that in Rana is connected with considerable chromosome rearrangements. The karyotype of this species differs from the karyotypes of many Rana species and from that of P. delalandii, 2n = 26, not only in the presence of the heteromorphic pair 8, but also in the presence of two pairs of telocentric chromosomes (10 and 11; Schmid, 1980a, b), which is very unusual for karyotypes of higher anurans (Morescalchi, 1973).

Relocation of NORs possibly also occurred in R. arvalis (with 2n = 24 and the secondary constrictions in the short arms of the 2nd pair – Ullerich, 1967) and R. dalmatina (with 2n = 26, the secondary constrictions are in the short arms of pair 3 – Guillemin, 1967; Popov, 1974). Chromosomes of these species have not been Ag-stained yet.

Besides the large NORs in the 10th pair, some North American and Asian Rana species have a few additional sites in different chromosomes which were positively stained by the Ag-AS-method (Schmid, 1978b; 1980a, b; Table 1). It is not yet clear if rDNA is localized in these additional sites, or if these Ag-positive regions are the sites of 5S DNA and other genes, as in the case of Triturus vulgaris meridionalis (Nardi et al., 1978). Moreover, in another urodelan species, Pleurodeles walt-

b The localization of NORs was confirmed by in situ hybridization of <sup>3</sup>H-18S-28S rRNA (Vitelli et al., 1982).

c NOR was localized only by in situ hybridization experiments (Vitelli et al., 1982).

d This pair is No. 11, but it is homologous to the pair No. 10 in the 26-chromosome Rana species (Green, 1983).

e This site appears to correspond to the 5S locus (Vitelli et al., 1982).

lii, silver preferentially stains the pericentromeric zones of chromosomes, where, after cold treatment of animals, additional secondary constrictions appear; the positions of these zones are matched by the positions of G-, C-, and Q-bands (Bailly, 1983). As for *R. catesbeiana*, it was shown by in situ hybridization of <sup>3</sup>H-18S + 28S rRNA and <sup>3</sup>H-5S rRNA that NORs are located only in the 10th pair, while the 5S DNA locus seems to be situated in Ag-positive sites of the 12th pair (Vitelly et al., 1982).

Unfortunately, the Ag-AS-method has not yet been used in many Rana species. Most interesting are the North American R. virgatipes (2n = 26), which has secondary constrictions in both arms of the homologues of the 10th pair (Green et al., 1981), and four Far Eastern Rana species, R. subaspera and R. holsi, 2n = 26, with secondary constrictions in two pairs, Nos. 4 and 9 (10) (Kuramoto, 1972), and R. namiyei and R. kuhlii, 2n = 22, with secondary constrictions in the homologues of the 6th and 14th pairs (Kuramoto, 1972; 1980).

As regards the additional secondary constrictions in the 24-chromosome species, *R. ornativentris* (pairs Nos. 2, 3, 4, and 7 – Seto, 1965), their positions are matched by heterochromatic C-bands without constrictions in *R. dybowskii* (Green, 1983). The position of the only secondary constriction in the short arms of *R. arvalis* chromosomes No. 2 (Ullerich, 1967) is also matched by the position of the C-band in *R. dybowskii* chromosome No. 2 (Green, 1983). As the NOR in *R. dybowskii* is associated with C-positive heterochromatin (Green, 1983), so the NOR in *R. arvalis* has probably been translocated into another C-heterochromatic region on the 2nd chromosome.

Our data and those shown in Table 1 point out that the localization of the NOR in Rana karyotypes is very conservative. Apparently less conservativeness was found in Bufo species: in Bufo karyotypes Ag-positive NORs may be located in at least three chromosome pairs (Schmid, 1978a; Beck & Mahan, 1979; Birstein, 1981a), In amphibians conservativeness of NOR localization may be characteristic of only some of the higher anurans, because even in the karyotypes of closely related species of Urodela (genera Triturus, Aneides, Plethodon), NORs are located in different chromosome pairs (Mancino et al., 1977; Macgregor & Sherwood, 1979).

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