Biometric data assessment on *Xenopus laevis* tadpoles

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Introduction

Recent studies stated that 41% of known species of amphibians are threatened.1 Indeed, their dramatic population decline is due to environmental conditions, including pollution.2,3 Their permeable skin and their close relationship with aquatic compartment, make them particularly sensitive to contamination and stand as strong models in biomonitoring studies.4 Consequently, the use of biomarkers in batrachians arose in the 80’s:5 from morphological and anatomical approaches (deformities, gonadic abnormalities) to biochemical measurements (vitellenogen, metallothionein, CYP450).4 For example, embryotoxic and teratogenic damages on larvae were detected with the FETAX (Frog Embryo Teratogenesis Assay - *Xenopus*) test developed in *Xenopus laevis*.6-8 In this regard, this specie *Xenopus laevis* was proposed regarding development stage, segmentation produces binary information such as spine integrity or brain appropriately developed regarding development stage, segmentation provides quantitative measurement of tadpole morphology. In this study, we focused on the tadpole’s length (calculated based on the Feret’s diameter, corresponding to the longest distance between two points of the selection boundary), perimeter, distance between eyes and position of the suction cup. Parameters can also be combined, normalizing measurements on animal and/or reference size to assess relative morphological differences.

Results and Discussion

Our new method provides numerous qualitative and quantitative data could be obtained. As depicted in Figure 2, a control tadpole (NT) was compared to treated ones (A, B and C). Morphological observations revealed abnormalities in terms of growth (tadpole A), dorsal curvature (tadpole B), or dorsal break and various edema (tadpole C).

While traditional approaches focus only on growth, mortality and qualitative data, our method relies on extensive images enhancement automation and complete this qualitative approach with new biometric criteria, and thus for each suited tadpole. Figure 2D shows some of these measurements: length, perimeter, distance between the eyes (eyes) and the position of the suction cup.

To go even farther in the approach, in a second step of experiment, different tadpoles were compared. First normalization (grey bars, Figure 2D) was done in comparison with a control organism (yellow dotted line, Figure 2D). Then, in order to highlight subtle alterations, a second normalization was achieved: each measurement was expressed as a function of studied tadpole length (colored bars). Thus, for treated tadpole A, it appeared that even if the individual was smaller (length, grey bar) the other measured parameters were over-size compared to untreated tadpole (colored bars).

Conclusions

The new method we propose presents several advantages compared to the traditional ones. Indeed, automated segmentation and detection allow automated quantification of tadpole’s morphology and anatomy. In this paper, we focused on several parameters such as perimeter or distance between eyes, but numerous others can be obtained from segmented images (*i.e.*, branchial arches or brain size and morphology, etc.). Thus, from the high quality of each individual description, it answers to the ethical principles of the 3R’s, first stated by Russell and Burch in 1959,10 by decreasing the number of tadpoles needed to...
achieve precise estimation of xenobiotic impact. Then, each contaminant results in a specific biometric signature, an even more efficient biomarker. Thus, this biometric data assessment appears as an effective tool for environmental biomonitoring.

References

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