Manuscript draft

Title – Genome annotations for higher mammals, a new frontier or reproducing old knowledge?

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Abstract

The development of high-throughput sequencing techniques continues, but the challenges of annotating genomes remain. Collection of relevant data and merging the results is an ongoing task.

Keywords: Annotations, development, mammals

Introduction

As scientists we are infallible. Everything we do are well thought through and always produces relevant and positive results. So was the case in this study.

Materials and Methods

We did stuff. Quite good stuff, I believe.

Results

Plenty of results, but had to be tailored to be useful. Corroborates our ideas perfectly now.

Discussion

We did nothing wrong, we were merely creative with the little means we had. Data was tortured to confession, but in a good way.

References

To be filled in.