Application Note:
JSAV: JavaScript Sequence Alignment Viewer

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ABSTRACT

Summary: JSAV

Availability and Implementation: JSAV is implemented in JavaScript. The code is documented using JSDOC and is available from `http://www.bioinf.org.uk/software/jsav/` or from GitHub `http://www.github.com/AndrewCRMartin/JSAV/`

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1 INTRODUCTION

2 IMPLEMENTATION

3 AVAILABILITY

The software may be downloaded from `http://www.bioinf.org.uk/software/jsav/` or from `http://www.github.com/AndrewCRMartin/JSAV/` and used freely. The web site provides demonstrations and full documentation.

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