

# The SeqWare Framework: Infrastructure Tools for Genomics

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## Abstract

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**Background:** Text for this section of the abstract.

**Results:** Text for this section of the abstract ...

**Conclusions:** Text for this section of the abstract ...

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## Background

Text for this section. [1–13]

## Results and Discussion

### Results sub-heading

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Sub-sub-sub-headings are made with the

*subsubsection* command.

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Text for this sub-sub-section ...

*Another sub-sub-sub-heading*

Text for this sub-sub-section ...

### **Another results sub-heading**

Text for this sub-section ...

### **Yet another results sub-heading**

Text for this sub-section. More results ...

## **Conclusions**

Text for this section ...

## **Methods**

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### **Yet another sub-heading for this section**

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## **Authors contributions**

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## **Acknowledgements**

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## **References**

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## Figures

### Figure 1 - Sample figure title

A short description of the figure content should go here.

### Figure 2 - Sample figure title

Figure legend text.

## Tables

### Table 1 - Sample table title

Here is an example of a *small* table in L<sup>A</sup>T<sub>E</sub>X using `\tabular{...}`. This is where the description of the table should go.

My Table		
A1	B2	C3
A2	...	..
A3	..	.

### Table 2 - Sample table title

Large tables are attached as separate files but should still be described here.

## **Additional Files**

### **Additional file 1 — Sample additional file title**

Additional file descriptions text (including details of how to view the file, if it is in a non-standard format or the file extension). This might refer to a multi-page table or a figure.

### **Additional file 2 — Sample additional file title**

Additional file descriptions text.