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Nehil Jain <sup>1,\*</sup>, Kacper Zukowski <sup>2</sup> and Co-Author <sup>2\*</sup>

<sup>1</sup>Department of Computer Science, Address XXXX etc.

<sup>2</sup>Department of XXXXXXXX, Address XXXX etc.

Received on XXXXX; revised on XXXXX; accepted on XXXXX

Associate Editor: XXXXXXX

#### **ABSTRACT**

**Motivation:** Many tools for GWAS and NGS in Bioconductor. Very few tools for pooled seq gwas in R. large datasets in R because of new hig performance packages. Works well on desktop and cluster configurations. Open source. Fast and intuitive methods for biology scientist to perform gwas analysis.

Contact: name@bio.com

### 1 INTRODUCTION

$$\sum x + y = Z \tag{1}$$

## 2 APPROACH

# 3 METHODS

 

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<sup>\*</sup>to whom correspondence should be addressed

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row2	row2	row2	row2
row3	row3	row3	row3
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Fig. 2. Caption, caption.

### 4 DISCUSSION

### 5 CONCLUSION

 

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### **ACKNOWLEDGEMENT**

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Funding: Text Text Text Text Text Text Text.

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