Using Biomedical Natural Language Processing to Identify Related Datasets

This manuscript (<u>permalink</u>) was automatically generated from <u>J-Wengler/NLP_Paper@9d1b117</u> on November 12, 2020.

Authors

• Stephen Piccolo

· 🕥 <u>srp33</u>

College of Life Science, Brigham Young University

• James Wengler

· 🕥 <u>J-Wengler</u>

College of Life Science, Brigham Young University

Abstract

Introduction

Methods

Idea

Testing markdown script

Results

Discussion

References