## **Project 9: Structural Health Monitoring**

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## 1. Introduction

## 1.1 Motivation behind the research experiment

Identification of damage from the analysis of vibration signals has received significant attention in the civil, mechanical and aerospace fields. Structural health monitoring allows the engineer to use sensing of the structural responses in conjunction with appropriate analysis and modeling techniques, to monitor the condition of a structure. The problem most commonly considered is that where data is recorded at two different times and it is of interest to determine if the structure suffered damage in the time interval between the two observations. The behavior of the system during the observation periods is typically assumed linear and the damage is identified as changes in system parameters. A solution can be obtained in principle by using the measured data to optimize a model of the structure in the two states and inspecting the differences. The goal of the experiment was to develope an automated structural health monitoring system capable of providing early warnings against structural damage. In order to achieve this, damage was simulated by removing bracing within the structure in nine different ways as shown in the following sections.

### 1.2 Literature Review

Due to the importance of this research, there have been multiple papers regarding how to optimize the recollection of the data and quantity needed for training the machine learning algorithm. Examination of the literature reveals, however, that the assumptions used to establish the various approaches vary widely and it is unclear what is the true capability of the current state of the art in damage detection of civil engineering structures.

# 1-D CNNs for structural damage detection: Verification on a structural health monitoring benchmark data

Nowadays, a large number of measurement scenarios are needed to generate training data before placing the sensors in large civil structures. The paper "1-D CNNs for structural damage detection: Verification on a structural health monitoring benchmark data" provides an enhanced CNN-based algorithm which only requires two training sessions, ensuring accuracy and speed. The damage was estimated successfully by applying a CNN-based approach only requiring two measurement datasets. The overall structural health monitoring procedure for this paper took place on a very short amount of time, around 5000x faster than the conventional procedure. The authors indicated that this approach can be utilized on any structures, regardless of the size of the structure. Yet, the data processed was based exclusively on a monitored structure, and one may wonder how the fully damaged scenario data could be obtained on a large civil structure that needs to be monitored. Also, other difficulties become more apparent if we take into consideration how large civil structures do not behave precisely as the replica in which accelerometers were placed. It is almost impossible to repeat the measurement procedure that took place for the Phase II benchmark study on any other structure. And finally, although anomaly detection is obtained using CNNs algorithms, the cause of such anomaly or the location is completely a mystery. It is important to know the state of a structure for public safety, but it is also very important to locate the damage in order to take action.

## 1.3 Structure & experiment descriptions:

#### **Benchmark Structure**

The benchmark structure is a 2-bay by 2-bay, 4 story steel frame structure at the University of British Columbia. Especially Benchmark Structure. Loaded from the latest version of image on GitHub.

Cases with known and unknown input and damage scenarios including symmetrical and unsymmetrical loss of stiffness in the bracing system were considered. The experiment in question is regarding how damage can be simulated by removing bracing or loosening bolts within a four-story steel frame structure. Complete details of the damage cases, input excitation and other pertinent aspects of the study of phase I can be found below. To obtain the data, accelerometers were placed throughout the structure to provide measurements of the structural responses. In particular, three sensors per floor. One located at the center, one at the west side and one at the east side, as the MATLAB files indicate. Then, different cases took place in which members were loosen or removed to analyze the output and correlate the difference in acceleration values with the difference in setup.

**The different cases** \* Case 1 - Fully braced configuration. \* Case 2 - All east side braces removed. \* Case 3 - Removed braces on all floors in one bay on southeast corner. \* Case 4 - Removed braces on 1st and 4th floors in one bay on southeast corner. \* Case 5 - Removed braces on 1st floor in one bay on southeast corner. \* Case 6 - Removed braces on all floors on east face, and 2nd floor braces on north faces. \* Case 7 - All braces removed on all faces. \* Case 8 - Configuration 7 + loosened bolts on all floors at both ends of beam on east face, north side. \* Case 9 - Configuration 7 + loosened bolts on floors 1 and 2 at both ends of beam on east face, north side.



**Excitation** Ambient vibration was inputted into the structure by two types of forced excitations. The forced exci- tation cases consider both impact hammer tests, and broadband excitations provided by an electrodynamic shaker. The choice of these two methods is to simulate the structure's response during an earthquake.

# 1.4 Motivation behind the project on structural damage detection

Numerous structural health monitoring algorithms have been developed and been implemented on experimental and full-scale structure. Because the techniques are applied to different structures under various conditions, the relative merits of each algorithm are not obvious. Thus, the community would benefit from a comparison of several algorithms when applied to the same problems.

## **Machine Learning Methods**

Methods go here

### **Results**

Results go here

## **Discussion of Results**

Discussion goes here

## Conclusion

Conclusion goes here

### References

#### 1. Sci-Hub provides access to nearly all scholarly literature

Daniel S Himmelstein, Ariel Rodriguez Romero, Jacob G Levernier, Thomas Anthony Munro, Stephen Reid McLaughlin, Bastian Greshake Tzovaras, Casey S Greene

eLife (2018-03-01) https://doi.org/ckcj

DOI: 10.7554/elife.32822 · PMID: 29424689 · PMCID: PMC5832410

#### 2. Reproducibility of computational workflows is automated using continuous analysis

Brett K Beaulieu-Jones, Casey S Greene

Nature biotechnology (2017-04) https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6103790/

DOI: <u>10.1038/nbt.3780</u> · PMID: <u>28288103</u> · PMCID: <u>PMC6103790</u>

#### 3. Bitcoin for the biological literature.

Douglas Heaven

Nature (2019-02) https://www.ncbi.nlm.nih.gov/pubmed/30718888

DOI: 10.1038/d41586-019-00447-9 · PMID: 30718888

## 4. Plan S: Accelerating the transition to full and immediate Open Access to scientific publications

cOAlition S

(2018-09-04) https://www.wikidata.org/wiki/Q56458321

#### 5. Open access

Peter Suber *MIT Press* (2012)

ISBN: <u>9780262517638</u>

#### 6. Open collaborative writing with Manubot

Daniel S. Himmelstein, Vincent Rubinetti, David R. Slochower, Dongbo Hu, Venkat S. Malladi, Casey S. Greene, Anthony Gitter

Manubot (2020-05-25) <a href="https://greenelab.github.io/meta-review/">https://greenelab.github.io/meta-review/</a>

#### 7. Opportunities and obstacles for deep learning in biology and medicine

Travers Ching, Daniel S. Himmelstein, Brett K. Beaulieu-Jones, Alexandr A. Kalinin, Brian T. Do, Gregory P. Way, Enrico Ferrero, Paul-Michael Agapow, Michael Zietz, Michael M. Hoffman, ... Casey S. Greene

Journal of The Royal Society Interface (2018-04-04) https://doi.org/gddkhn

DOI: <u>10.1098/rsif.2017.0387</u> · PMID: <u>29618526</u> · PMCID: <u>PMC5938574</u>

#### 8. Open collaborative writing with Manubot

Daniel S. Himmelstein, Vincent Rubinetti, David R. Slochower, Dongbo Hu, Venkat S. Malladi, Casey S. Greene, Anthony Gitter

PLOS Computational Biology (2019-06-24) https://doi.org/c7np

DOI: 10.1371/journal.pcbi.1007128 · PMID: 31233491 · PMCID: PMC6611653

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The rest of this document is a full list of formatting elements/features supported by Manubot. Compare the input (.md files in the /content directory) to the output you see below.

## **Basic formatting**

#### **Bold text**

Semi-bold text

Centered text

Right-aligned text

Italic text

Combined italics and bold

#### Strikethrough

- 1. Ordered list item
- 2. Ordered list item
  - a. Sub-item
  - b. Sub-item
    - i. Sub-sub-item
- 3. Ordered list item
  - a. Sub-item
- · List item
- · List item
- List item

subscript: H<sub>2</sub>O is a liquid

superscript: 2<sup>10</sup> is 1024.

unicode superscripts 0123456789

unicode subscripts 0123456789

A long paragraph of text. Lorem ipsum dolor sit amet, consectetur adipiscing elit, sed do eiusmod tempor incididunt ut labore et dolore magna aliqua. Ut enim ad minim veniam, quis nostrud exercitation ullamco laboris nisi ut aliquip ex ea commodo consequat. Duis aute irure dolor in reprehenderit in voluptate velit esse cillum dolore eu fugiat nulla pariatur. Excepteur sint occaecat cupidatat non proident, sunt in culpa qui officia deserunt mollit anim id est laborum.

Putting each sentence on its own line has numerous benefits with regard to <u>editing</u> and <u>version</u> <u>control</u>.

Line break without starting a new paragraph by putting two spaces at end of line.

## **Document organization**

Document section headings:

## **Heading 1**

## **Heading 2**

**Heading 3** 

Heading 4

Heading 5

Heading 6



#### Horizontal rule:

Heading 1's are recommended to be reserved for the title of the manuscript.

Heading 2's are recommended for broad sections such as Abstract, Methods, Conclusion, etc.

Heading 3's and Heading 4's are recommended for sub-sections.

### Links

Bare URL link: <a href="https://manubot.org">https://manubot.org</a>

<u>Long link with lots of words and stuff and junk and bleep and blah and stuff and other stuff and more stuff yeah</u>

Link with text

Link with hover text

Link by reference

### **Citations**

Citation by DOI [1].

Citation by PubMed Central ID [2].

Citation by PubMed ID [3].

Citation by Wikidata ID [4].

Citation by ISBN [5].

Citation by URL [6].

Citation by alias [7].

Multiple citations can be put inside the same set of brackets [1,5,7]. Manubot plugins provide easier, more convenient visualization of and navigation between citations [2,3,7,8].

Citation tags (i.e. aliases) can be defined in their own paragraphs using Markdown's reference link syntax:

## Referencing figures, tables, equations

Figure 1

Figure 2

```
Figure 3

Figure 4

Table 1

Equation 1

Equation 2
```

## **Quotes and code**

Quoted text

Quoted block of text

Two roads diverged in a wood, and I—I took the one less traveled by, And that has made all the difference.

Code in the middle of normal text, aka inline code.

Code block with Python syntax highlighting:

```
from manubot.cite.doi import expand_short_doi

def test_expand_short_doi():
    doi = expand_short_doi("10/c3bp")
    # a string too long to fit within page:
    assert doi == "10.25313/2524-2695-2018-3-vliyanie-enhansera-copia-i-
        insulyatora-gypsy-na-sintez-ernk-modifikatsii-hromatina-i-
        svyazyvanie-insulyatornyh-belkov-vtransfetsirovannyh-geneticheskih-
        konstruktsiyah"
```

Code block with no syntax highlighting:

```
Exporting HTML manuscript
Exporting DOCX manuscript
Exporting PDF manuscript
```

## **Figures**



**Figure 1:** A square image at actual size and with a bottom caption. Loaded from the latest version of image on GitHub.



**Figure 2:** An image too wide to fit within page at full size. Loaded from a specific (hashed) version of the image on GitHub.



Figure 3: A tall image with a specified height. Loaded from a specific (hashed) version of the image on GitHub.



**Figure 4:** A vector .svg image loaded from GitHub. The parameter sanitize=true is necessary to properly load SVGs hosted via GitHub URLs. White background specified to serve as a backdrop for transparent sections of the image.

## **Tables**

**Table 1:** A table with a top caption and specified relative column widths.

Bowling Scores	Jane	John	Alice	Bob
Game 1	150	187	210	105
Game 2	98	202	197	102
Game 3	123	180	238	134

**Table 2:** A table too wide to fit within page.

	Digits 1-33	Digits 34-66	Digits 67-99	Ref.
pi	3.14159265358979323 846264338327950	28841971693993751 0582097494459230	78164062862089986 2803482534211706	piday.org
е	2.71828182845904523 536028747135266	24977572470936999 5957496696762772	40766303535475945 7138217852516642	nasa.gov

 Table 3: A table with merged cells using the attributes plugin.

	Colors		
Size	Text Color	Background Color	
big	blue	orange	
small	black	white	

## **Equations**

A LaTeX equation:

$$\int_0^\infty e^{-x^2} dx = \frac{\sqrt{\pi}}{2} \tag{1}$$

An equation too long to fit within page:

$$x = a + b + c + d + e + f + g + h + i + j + k + l + m + n + o + p + q + r + s + t + u + v + w + x + y + z + 1 + 2 + 3 + 4 + 5 + 6 + 7 + 8 + 9$$
(2)

## **Special**

▲ WARNING The following features are only supported and intended for .html and .pdf exports. Journals are not likely to support them, and they may not display correctly when converted to other formats such as .docx .

LINK STYLED AS A BUTTON

Adding arbitrary HTML attributes to an element using Pandoc's attribute syntax:

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Adding arbitrary HTML attributes to an element with the Manubot attributes plugin (more flexible than Pandoc's method in terms of which elements you can add attributes to):

Manubot Manubo

Available background colors for text, images, code, banners, etc:

white lightgrey grey darkgrey black lightred lightyellow lightgreen lightblue lightpurple red orange yellow green blue purple

Using the Font Awesome icon set:



**Light Grey Banner** useful for *general information* - <u>manubot.org</u>

## **1** Blue Banner

useful for important information - manubot.org

**♦ Light Red Banner** useful for *warnings* - <u>manubot.org</u>