

Ecological Succession in Central Park after Catastrophic Occurrences Cristina Guzman-Moumtzis¹, Jatna Paulino¹, Alyssa Simon¹, Vincent Joralemin¹ ¹Frank McCourt High School

Abstract

Ecological Succession is the observable change of an ecosystem over time. Central Park undergoes a type of artificial ecological succession but their impact is uncertain. Using a standard DNA extraction method, fifteen species of plants were studied to determine the impact of ecological succession in Central Park. A majority of the plants were nonnative and had a negative impact but some of the plant's were native and even endangered. This increase in diversity suggests that while ecological succession does allow for invasive species to enter an area, it also allows for weaker native species to thrive.

Introduction

In 1979, Wayne P. Sousa investigated changes to a specie of algae once the original species was devastated and a pioneer specie of algae began to colonize the area. The species of algae changed repeatedly until it it settled on one dominant algae¹ but before then, the ecosystem fluctuated dangerously. In comparison, those who observe the forest fires in Yellowstone Park claim these events are a key factor in shaping the ecology of the park.² Most positive examples, such as that of Yellowstone Park, is said to be those areas with semi-frequent events of ecological succession (according to the Intermediate Disturbance Hypothesis³). If species are too frequently replaced, the ecosystem will not have the time to adapt but if no change occurs, diversity decreases.

Methods

The samples for this project were collected from a recently renovated area of Central Park;a construction site located on West 81st. Four locations around the construction were chosen as well as four control locations six steps away from the site.Afterwards, all plants which appear different will be collected from both sites to collect a good sample size, had their locations marked and later recorded onto a sample database. By using a standard DNA extraction method ⁴ the DNA from each of the samples was extracted, amplified and then barcoded.

Results

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Database	Species	Origins ⁵
Identifier		
	Plantago	
KEN-002	lanceolata	Nonnative
	Plantago	
KEN-003	lanceolata	Nonnative
KEN-004	Poa Annua	Nonnative
	Symphyotrichum	
KEN-006 F	ciliolatum	Native
	Symphyotrichum	
KEN-008	laeve	Native
	Adenocaulon	
KEN-009	bicolor	Nonnative
KEN-010	Vicia ramiflora	Nonnative
KEN-011	Poa annua	Nonnative
	Plantago	
KEN-012	lanceolata	Nonnative
	Plantago	
KEN-014	lanceolata	Nonnative
KEN-015	Potentilla reptans	Nonnative
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Database	Potentilla reptans Species	Nonnative Origins ⁵
	Species	
Database Identifier	Species Lolium	Origins ⁵
Database Identifier KEN-030	Species Lolium temulentum	Origins ⁵ Nonnative
Database Identifier KEN-030 KEN-031	Species Lolium temulentum Trifolium repens	Origins ⁵ Nonnative Nonnative
Database Identifier KEN-030 KEN-031 KEN-032	Species Lolium temulentum Trifolium repens Plantago asiatica	Origins ⁵ Nonnative Nonnative Nonnative
Database Identifier KEN-030 KEN-031 KEN-032 KEN-033	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americana	Origins ⁵ Nonnative Nonnative Nonnative Native
Database Identifier KEN-030 KEN-031 KEN-032 KEN-033 KEN-034	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americanaTrifolium repens	Origins ⁵ Nonnative Nonnative Nonnative Native Nonnative
Database Identifier KEN-030 KEN-031 KEN-032 KEN-033	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americanaTrifolium repensTrifolium repens	Origins ⁵ Nonnative Nonnative Nonnative Native
Database Identifier KEN-030 KEN-031 KEN-032 KEN-033 KEN-034 KEN-035	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americanaTrifolium repensTrifolium repensLolium	Origins ⁵ Nonnative Nonnative Nonnative Nonnative Nonnative
Database Identifier KEN-030 KEN-031 KEN-032 KEN-033 KEN-034 KEN-035	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americanaTrifolium repensTrifolium repensLolium temulentum	Origins ⁵ Nonnative Nonnative Nonnative Nonnative Nonnative
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Database Identifier KEN-030 KEN-031 KEN-032 KEN-033 KEN-034 KEN-035	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americanaTrifolium repensTrifolium repensLolium temulentumPlantago asiaticaTrifolium repensTrifolium repensTrifolium repensTrifolium repensTrifolium repensTrifolium repensItemulentumItemulentu	Origins ⁵ Nonnative Nonnative Nonnative Nonnative Nonnative
Database Identifier KEN-030 KEN-031 KEN-032 KEN-033 KEN-034 KEN-035 KEN-036 KEN-037	SpeciesLolium temulentumTrifolium repensPlantago asiaticaUlmus americanaTrifolium repensTrifolium repensLolium temulentumPlantago asiaticaTrifolium repensTrifolium repensTrifolium repensTrifolium repensTrifolium repensTrifolium repensTrifolium repensTrifolium temulentumPlantago asiaticaTrifolium montanumTrifolium montanum	Origins ⁵ Nonnative Nonnative Nonnative Nonnative Nonnative Nonnative Nonnative Nonnative Nonnative
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This table represents all of the barcoded DNA collected through this procedure. The samples begin at KEN-002 and all unsuccessfully extracted DNA is omitted from the table. In addition to the species name and database identifier, the table also includes a column labeling each plant as a native specie or a non native specie (a specie which was introduced to the country).

Database Identifier	Species	Origins ⁵
KEN-016	Potentilla reptans	Nonnative
KEN-017	Plantago asiatica	Nonnative
KEN-018	Adenocaulon bicolor	Nonnative
KEN-019	Plantago lanceolata	Nonnative
KEN-020	Poa pratensis	Native
KEN-021	Trifolium montanum	Nonnative
KEN-022	Plantago asiatica	Nonnative
KEN-023	Trifolium montanum	Nonnative
KEN-024	Lolium temulentum	Nonnative
KEN-028	Potentilla reptans	Nonnative
KEN-029	Poa supina	Nonnative
Database Identifier	Species	Origins ⁵
KEN-041	Trifolium montanum	Nonnative
KEN-042	Plantago asiatica	Nonnative
KEN-043	Poa supina	Nonnative
KEN-044	Trifolium montanum	Nonnative
KEN-045	Poa supina	Nonnative
KEN-046	Prunus spinosa	Nonnative
KEN-048	Prunus spinosa	Nonnative
KEN-049	Poa pratensis	Native
KEN-051	Anthoxanthum nitens	Native
KEN-052	Plantago asiatica	Nonnative
KEN-053	Plantago lanceolata -	Nonnative

Central Park's plant diversity ranges from indigenous and introduced species⁶ but areas without disruptive events have lower biodiversity. This may be attributed to the lack of care that area of construction have in terms of chemicals such as RoundUp⁷, which prevents protein production in plants that utilize EPSP synthase⁸. While this herbicide does remove non native species of plants such as dandelions, it can has a major effect on other native plants. The genetic diversity of areas sprayed with this herbicide may hinder the growth of non native and native species but more research will need to be conducted for a definitive answer.

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⁵"Welcome to the PLANTS Database | USDA PLANTS." *Welcome to the PLANTS* Database | USDA PLANTS. Web. 20 May 2016. < http://plants.usda.gov/>.

⁶"Flora of Central Park." *Flora of Central Park*. Web. 27 May 2016. < http://www. nybg.org/files/scientists/datha/CentralPark/Maps.html>.

⁷Pang, Amelia. "Are NYC Parks Becoming More Toxic?" *The Epoch Times Are* NYC Parks Becoming More Toxic Comments. The Epoch Times, 18 Jan. 2016. Web. 27 May 2016. < http://www.theepochtimes.com/n3/1940550-are-nyc-parksbecoming-more-toxic/>.

⁸ "What Is Biology Good For? Controlling Weeds: RoundUp." *IUPUI Department* of Biology. Web. 25 May 2016. < http://www.biology.iupui. edu/biocourses/N100/goodfor13.html>.

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¹ Wayne P. Sousa (Last edited date). Experimental Investigations of Disturbance and Ecological Succession in a Rocky Intertidal Algal Community. <<u>http://www.</u> esajournals.org/doi/abs/10.2307/1942484>

² National Park Service U.S. Department of the Interior (2015). *General Format* Retrieved from http://www.nps.gov/yell/planyourvisit/upload/RI 2015 fire sm.

³ MICHAEL ALLABY. "intermediate disturbance hypothesis." A Dictionary of Ecology. 2004. *Encyclopedia.com*. 27 May. 2016 < http://www.encyclopedia.com>

⁴"Using DNA Barcodes to Identify and Classify Living Things." *Experiment* Protocol | DNA Barcoding 101. Cold Spring Harbor's Laboratory. Web. 15 Nov.

Acknowledgements