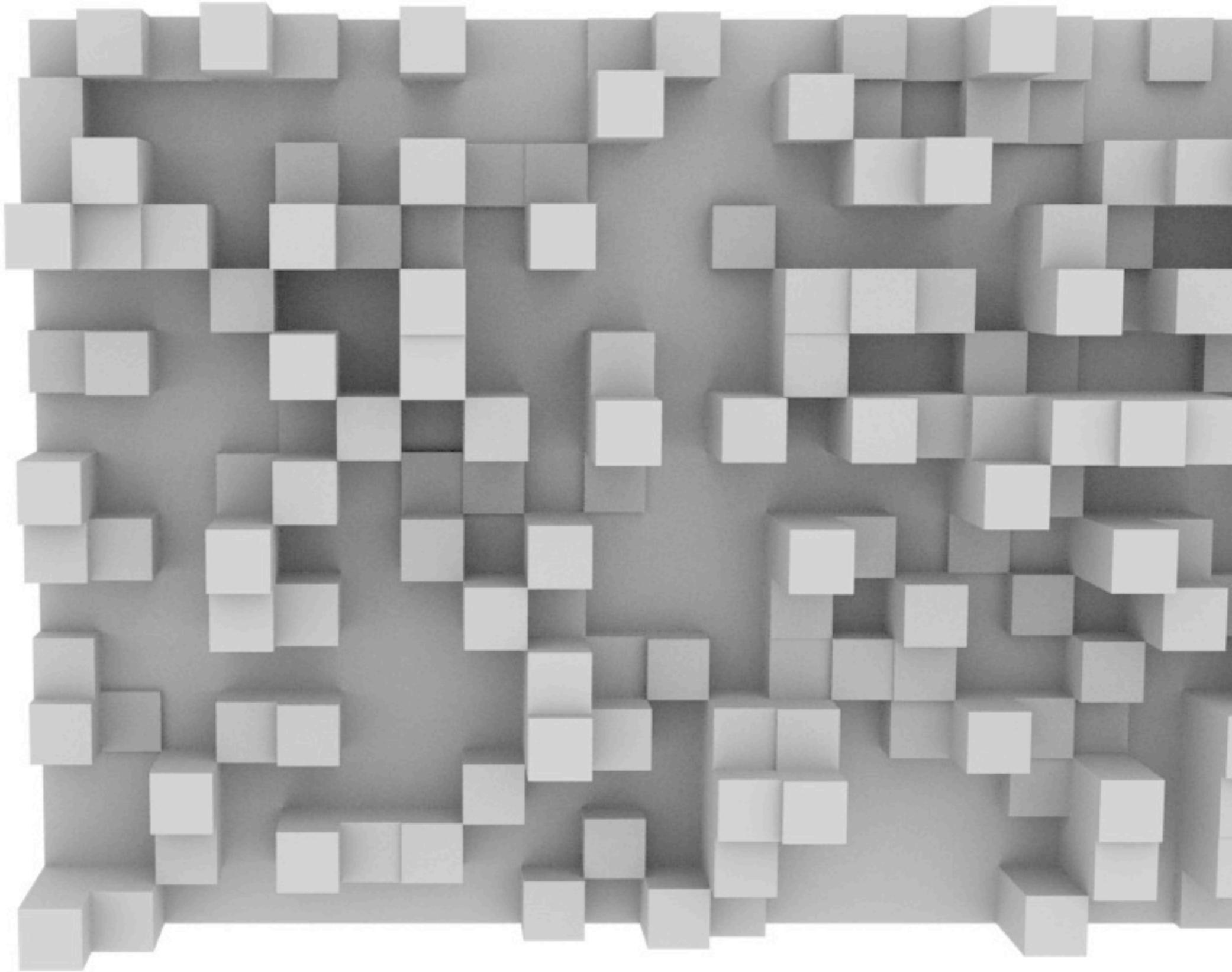


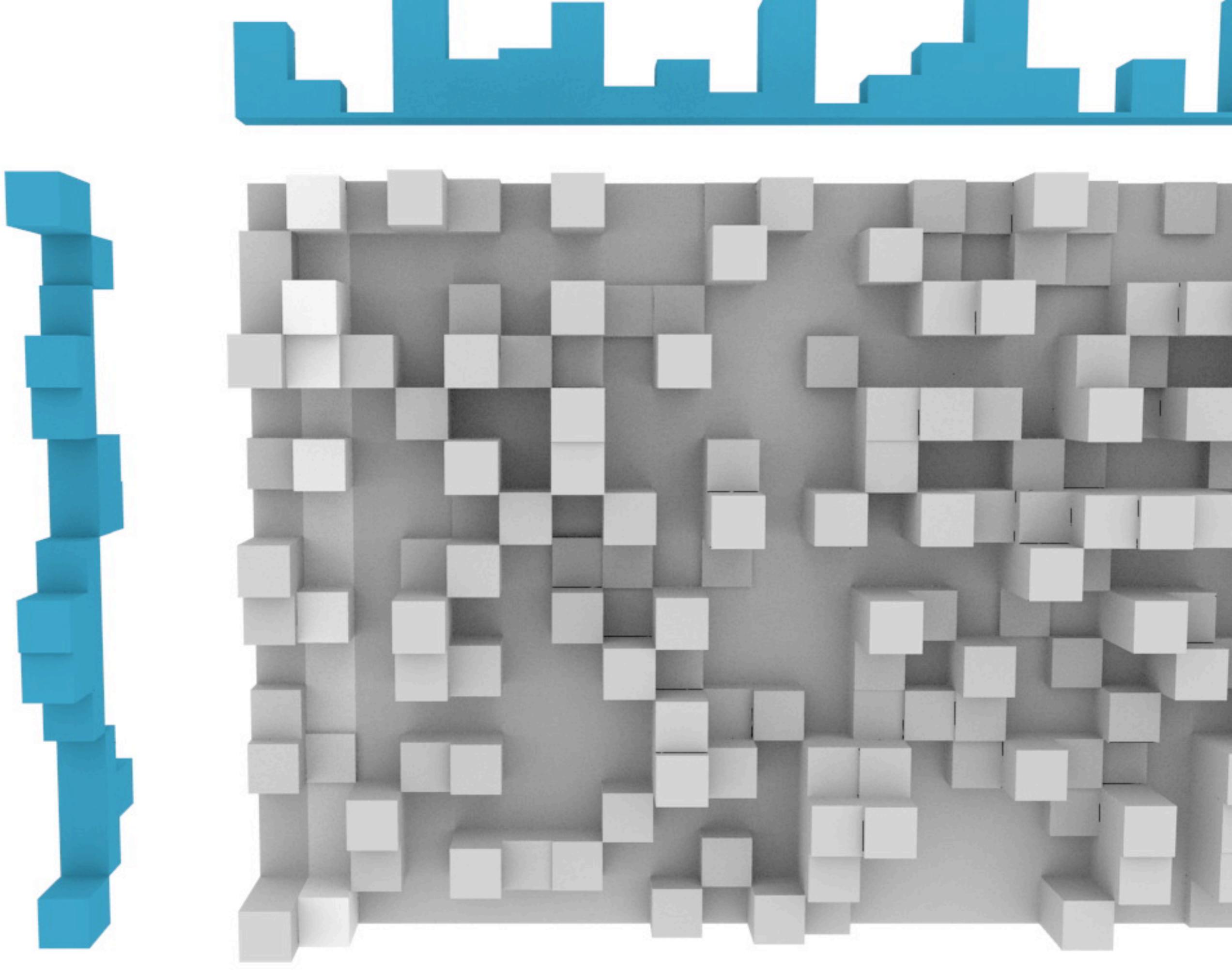
Signature



Treatments

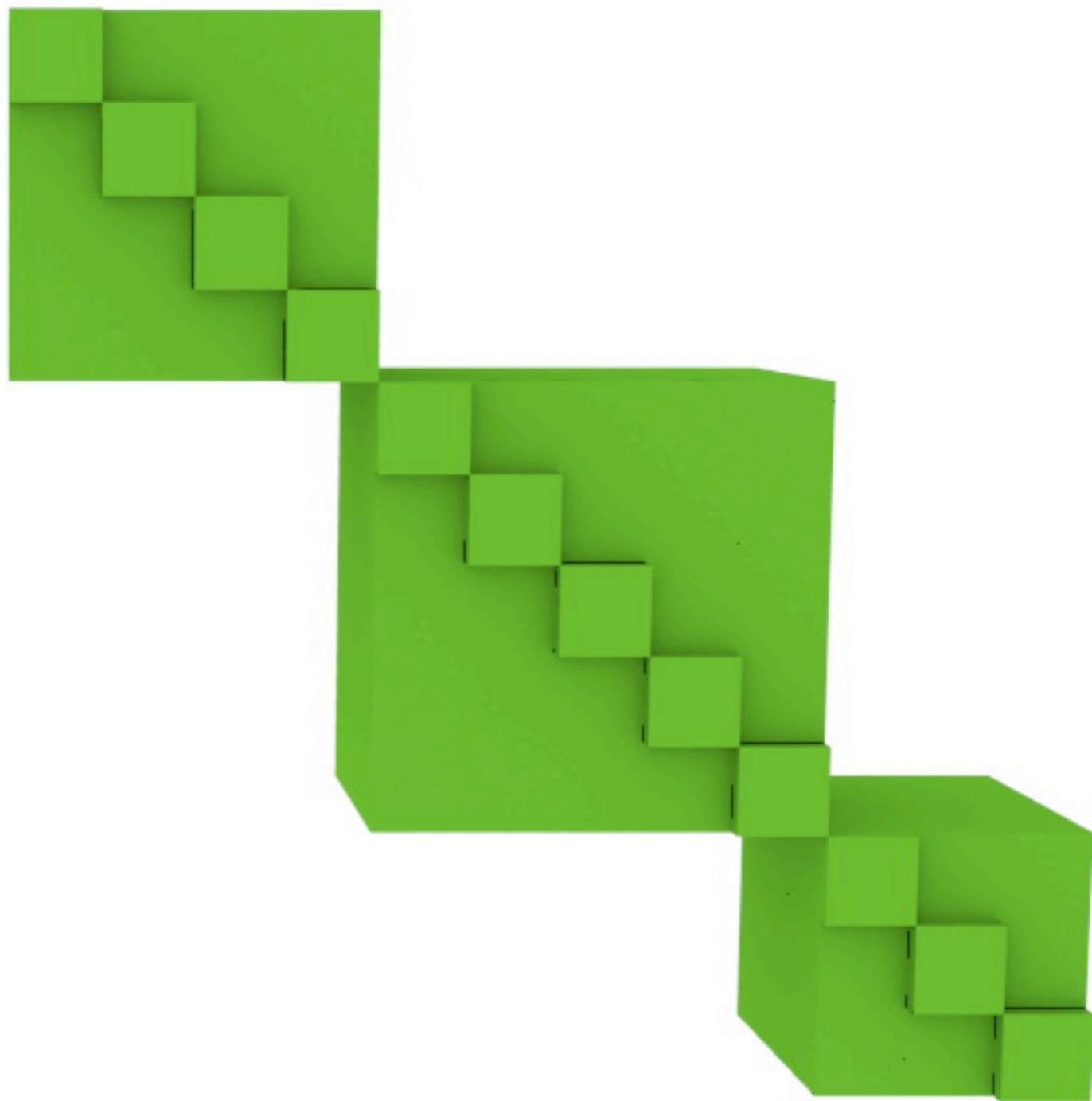
Genes





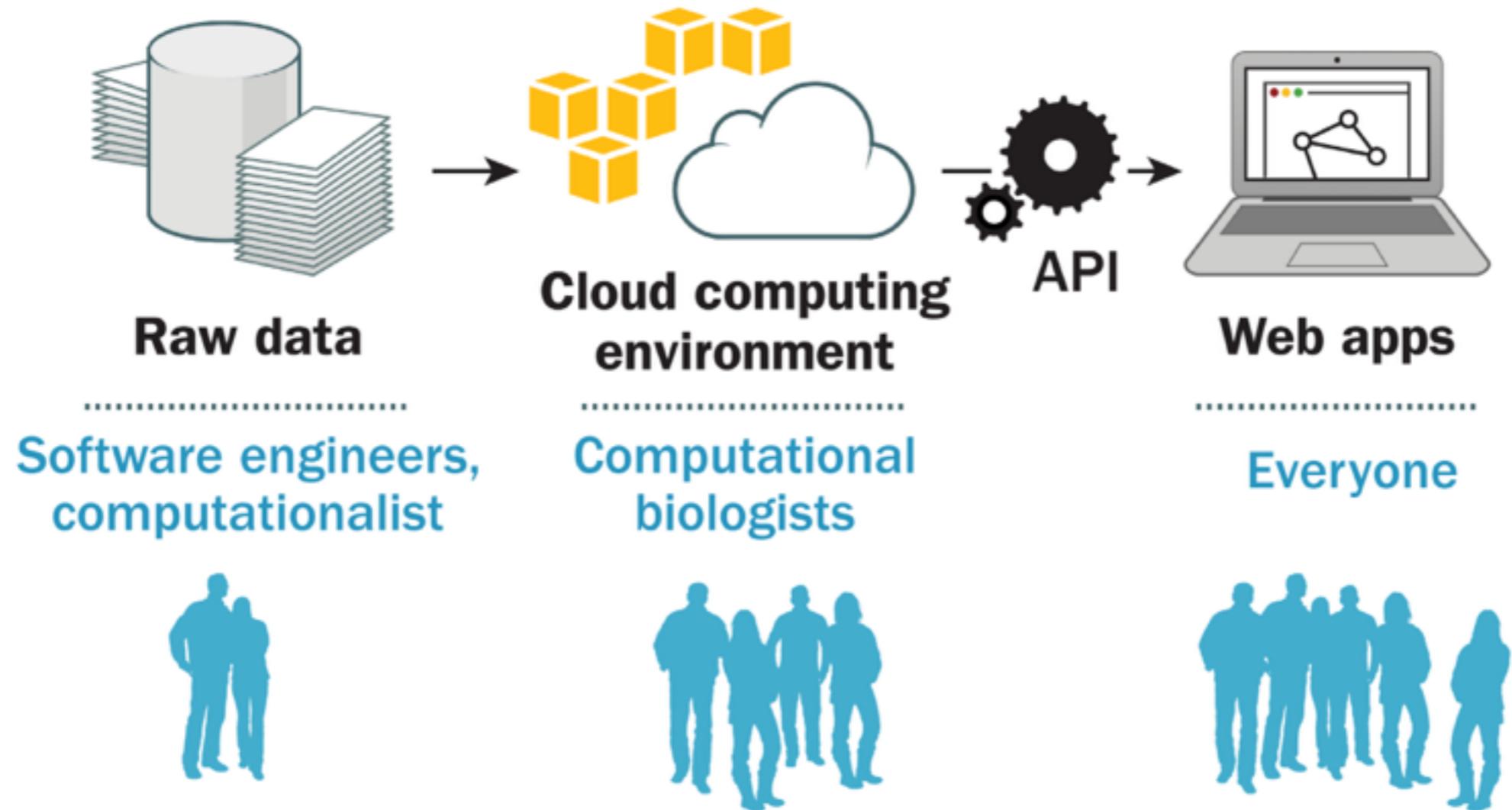
Treatments

Treatments

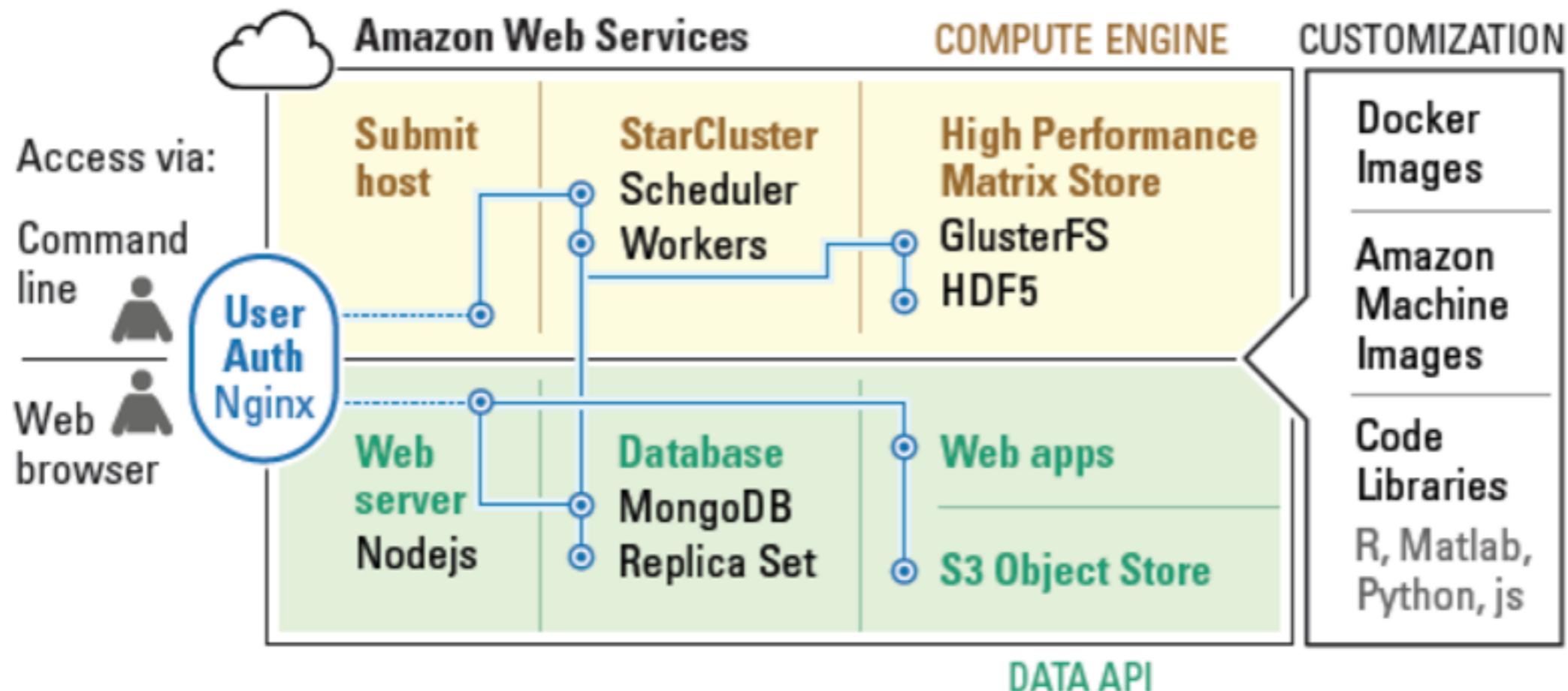


CLUE

Connectivity Map and L1000 User Environment



technology stack



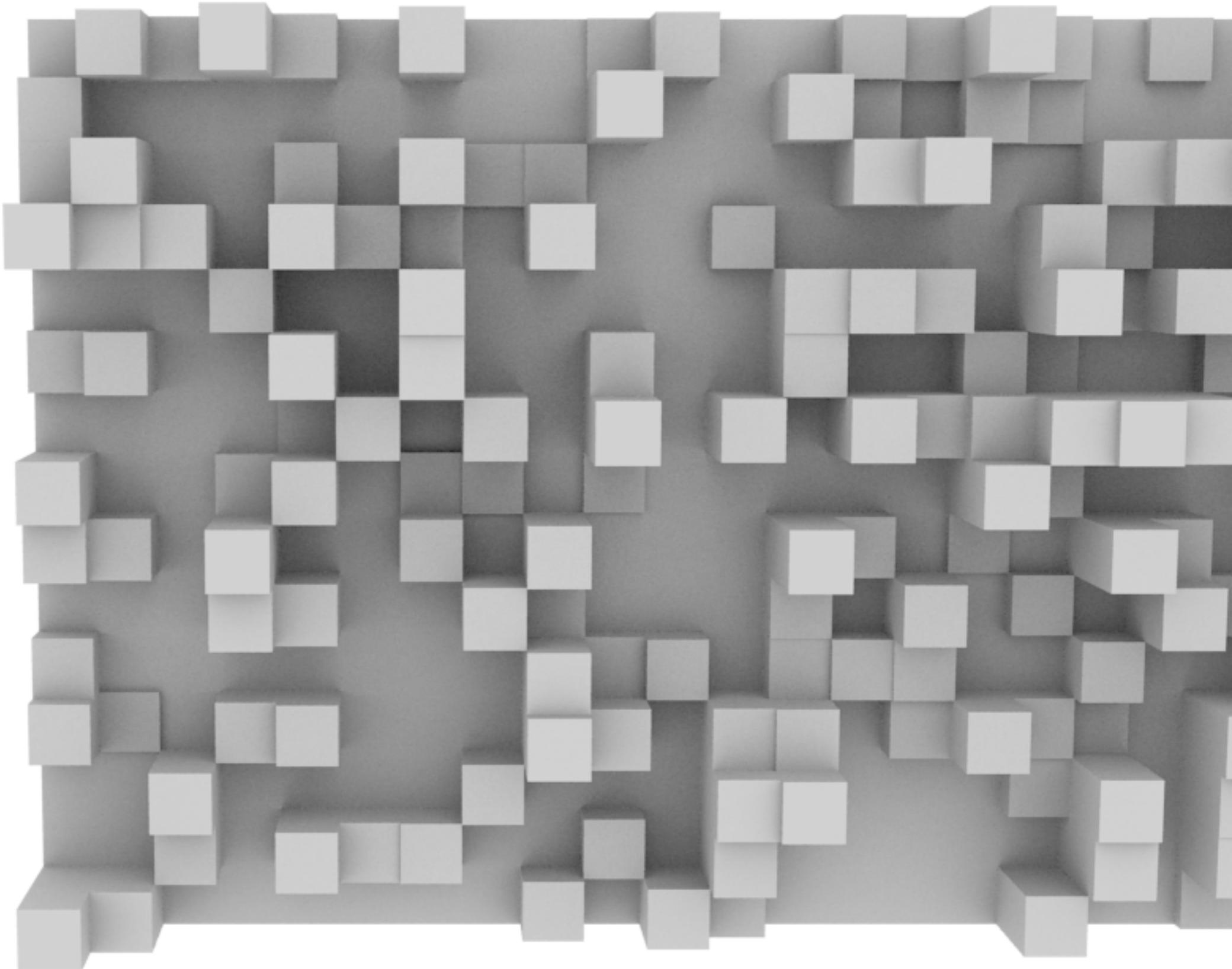


Apps





Treatments



[show an example](#)[take a tour](#)

GENETIC REAGENTS

0

knock down	0
over expression	0
variant png	0

CELLULAR CONTEXTS

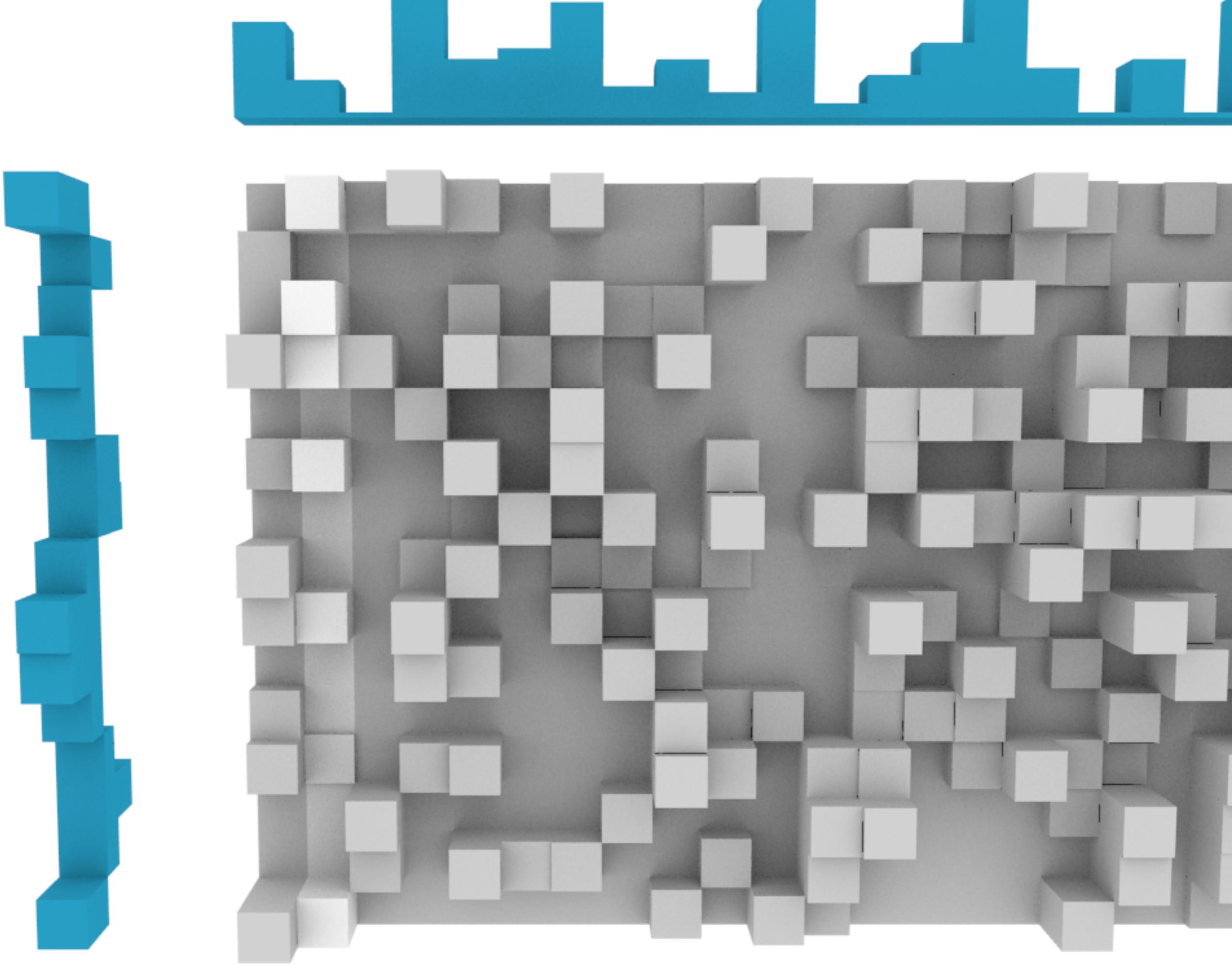
0

cancer	0
primary	0
other png	0

CHEMICAL REAGENTS

0

tool compounds	0
drugs and bioactives	0
other png	0

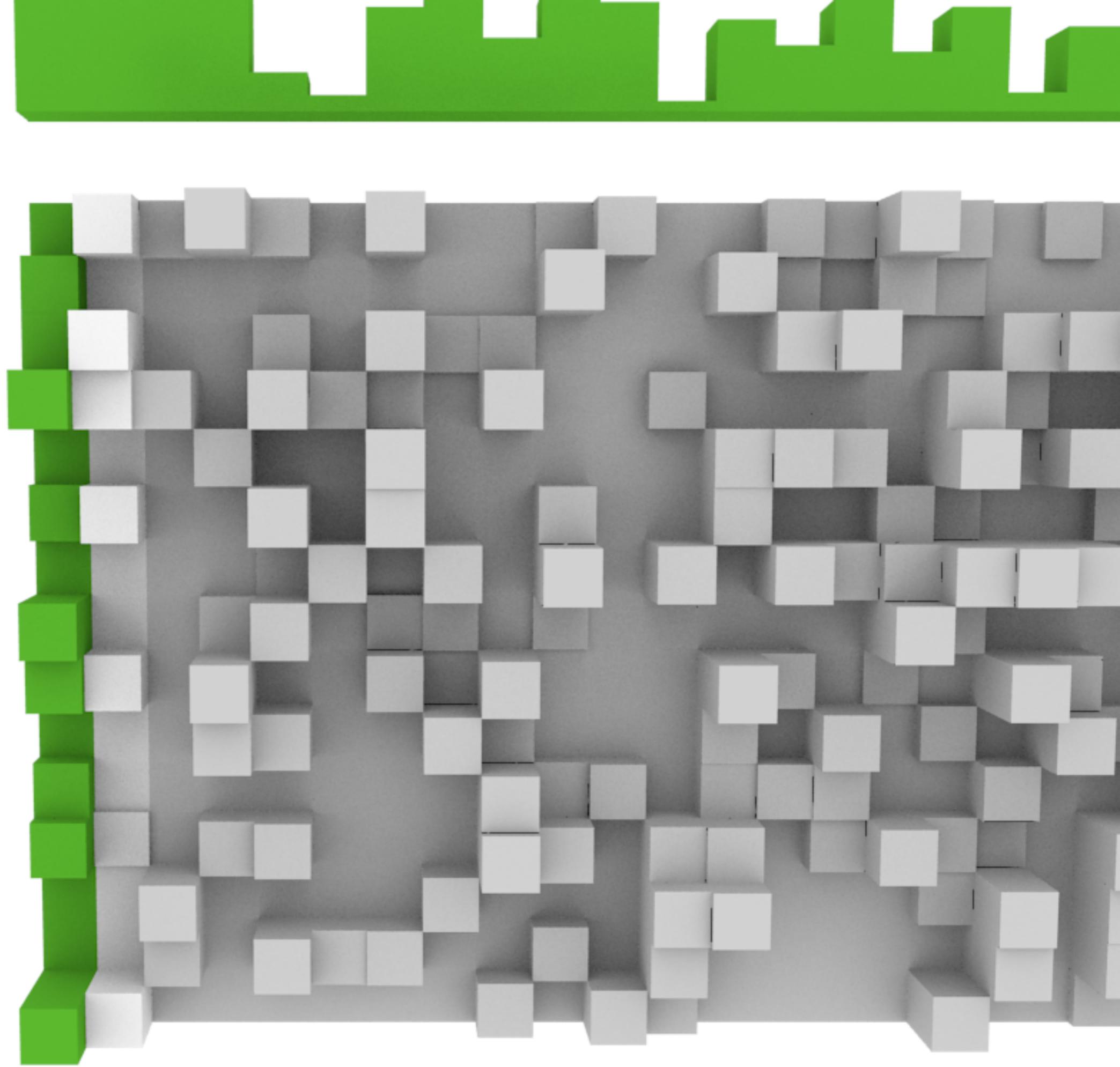


[take a tour](#)Name your query
➊ Enter Up-regulated genes

Enter one gene symbol or Affymetrix U133A probe ID per line or drag and drop a plain text file here.

➋ Enter Down-regulated genes

Enter one gene symbol or Affymetrix U133A probe ID per line or drag and drop a plain text file here.





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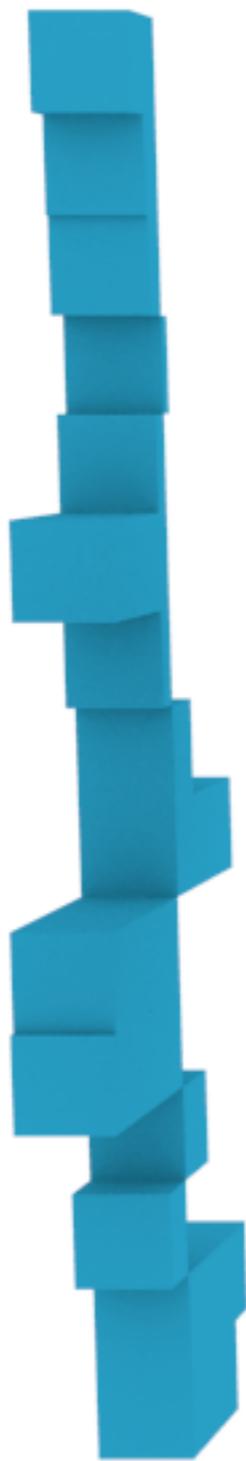
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CONNECTIVITY MAP

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Classes



0.1



0.9



0.3





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 CONNECTIVITY MAP

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API

[Overview](#)
[Usage](#)
[CellInfo](#)
[GeneInfo](#)
[InstInfo](#)
[PertInfo](#)
[PlateInfo](#)
[SigInfo](#)

Overview

The Lincscloud API offers programmatic access to annotations and perturbational signatures in [the LINCS L1000 dataset](#) via a collection of HTTP-based RESTful web services. These services support complex queries via simple HTTP GET requests that can be executed in a web browser or any programming language. The results are returned as standard JSON objects. Click on the links on the left for usage instructions and examples.

Live Demo

Click on the links on the left for API usage instructions and examples.

Access



Thank you for your interest in the LINCS Connectivity Map project. To learn more about the project and request access please visit lincscloud.org



Cloud

wmd91-702 [~]:

]

clue.io

free for academic users

7 day trial for private users