

# ***Web-Lar***

## ***First project results***

# ***Task assignments***

Extension of the LAR.js  
and LAR-DEMO.js  
projects with convex  
cells

- Andrea Somma
- Elisa Lamberti
- Fabio Cumbo

Integration of the matrix  
computation web-  
service

- Fabrizio Rebecca
- Luca Menichetti (**Group Leader**)

Integration of PDB  
visualization service

- Oscar Eijsermans

# Extension of the LAR.js and LAR- DEMO.js projects with convex cells

**Andrea Somma** assignment: 1-A, 1-B

**Elisa Lamberti and Fabio Cumbo** assignment: 2-A, 2-B



# Extension of the LAR.js and LAR-DEMO.js projects with convex cells

## Description

*Extension of the LAR.js and LAR\_DEMO.js projects with convex cells*

## Summary

### **1. Implementation of the cartesian product function between convex cell**

- Integration of the lar.js project with the **larProduct** function, which will allow to do the product between convex cells

### **2. Reimplementation of the "extract" function, generalizing it to make it work on cells with general shape**

- Implementation of the **larFacets** function in the lar.js project, currently provided with an extraction function that works only on simplexes.

# Extension of the LAR.js and LAR-DEMO.js projects with convex cells

## Brief explanation

*The lar.js wasn't provided of:*

- Cartesian product between convex cells
- Extract function

## Solution

So, the actual workflow for points 1 & 2 for today's demo was to:

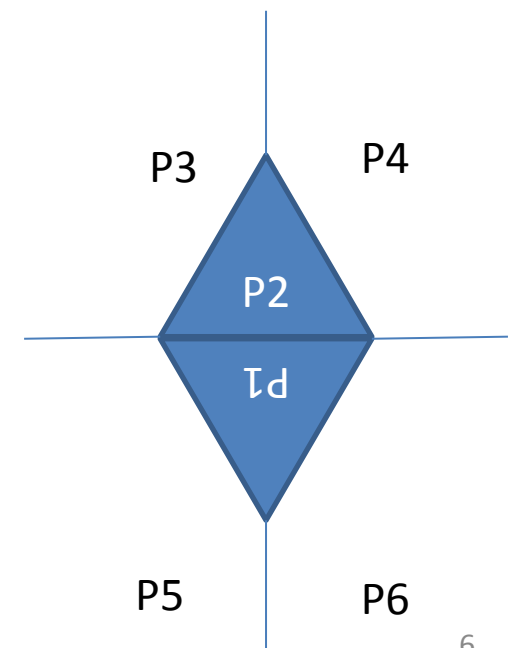
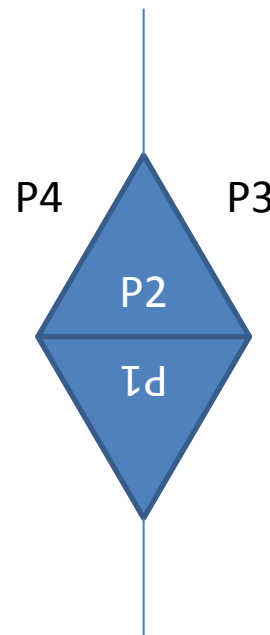
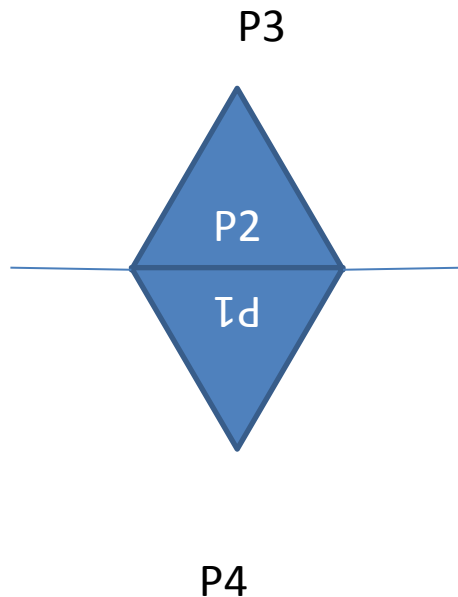
- ✓ Implement the lar.py functions "larProduct" & "larFacets" using dense matrixes, in lar.js.

Actually, we have readapted those functions and we have implemented a "beta version", which works well with simple models.

# Extension of the LAR.js and LAR-DEMO.js projects with convex cells

## To Do

- Other test and examples for Cartesian product
- **External space partitioning for larFacets:**  
**How to do that? What the criteria?**



# Extension of the LAR.js and LAR-DEMO.js projects with convex cells



# Integration of the matrix computation web-service

**Fabrizio Rebecca** assignment: communication  
**Luca Menichetti** assignment: handler





# Integration of the matrix computation web-service

## Description

*Create a webservice that provides fast matrix operations on the network*

## Summary

### **1. Localization of code's sections that use matrix operations**

- Replacement with a call to an external procedure (\*interface\*)

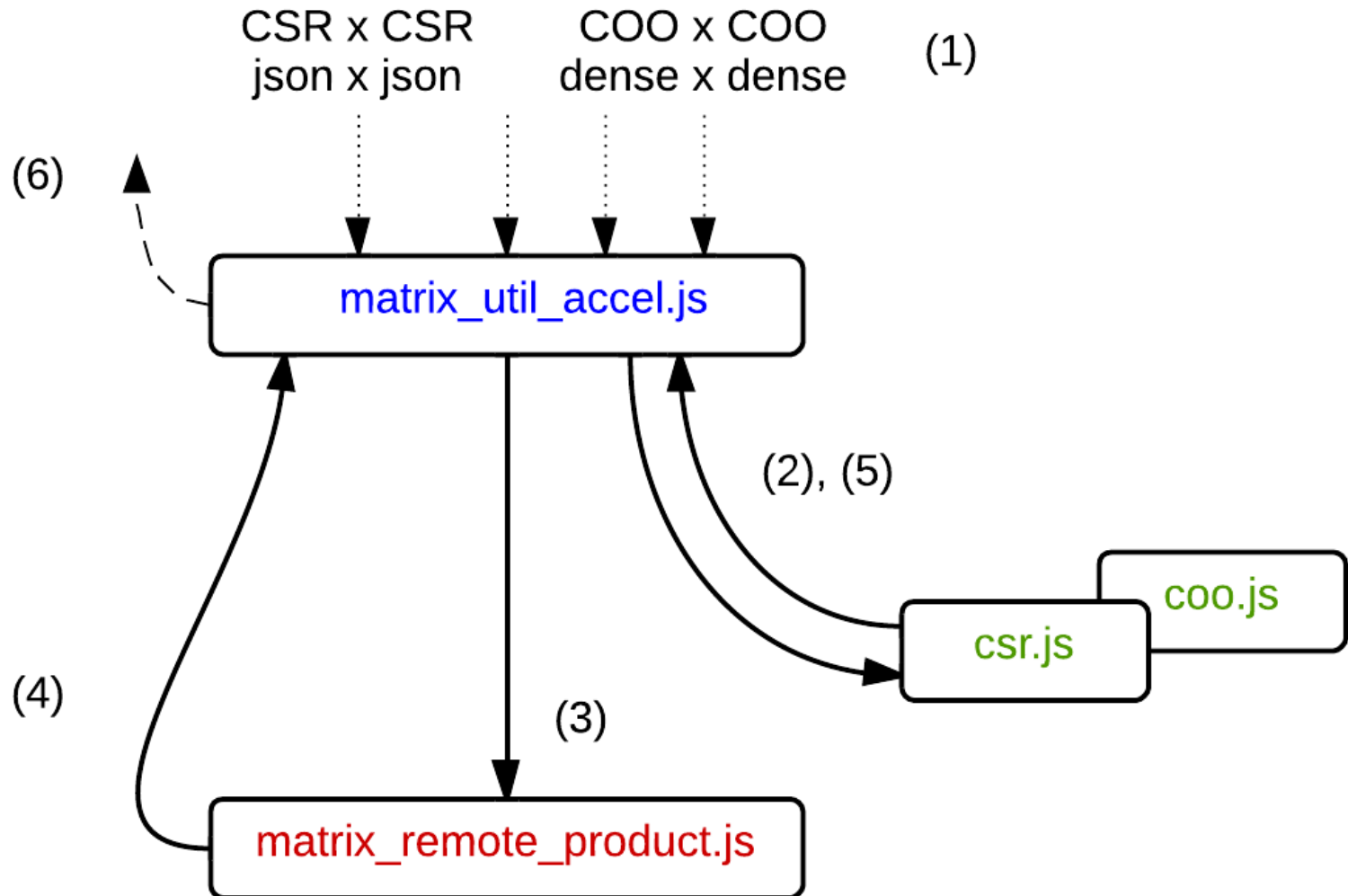
### **2. Creation of a proper layer that implements the computation of matrix's operation hiding the logic implementation**

- Definition of a **matrix\_util\_accel.js** layer (responsible to handle format or encoding, such as csr, coo, json)
- Realization of a HttpRequest with the matrix computation webservice using Representational state transfer – REST – in the **matrix\_remote\_product.js** layer (This layer is also responsible to manage the answer, with an opportune decoding in order to satisfy the specifics)

### **3. Setup a web service that offers such operation online**

- available with node.js;
- acquisition of the request using REST;
- executing the computations (OpenCL) and forwarding the results;
- create a *domain name* in which will be available such services online

# Integration of the matrix computation web-service



# Integration of the matrix computation web-service



# Integration of PDB visualization service

Oscar Eijsermans assignment



# Integration of PDB visualization service

## Description

*Integration of WebMOL and WebPDB projects*

## Summary

- 1. Identify the structure and functionalities of WebPDB and WebMOL**
  - Data structure and communication protocols
- 2. Identify the connections between the two projects**
  - Make changes where necessary
- 3. Merge the two projects in one project**

# Integration of PDB visualization service

## WebMOL-WebPDB interaction:

- **step 1** - install CouchDB and Node.js
- **step 2** - install node modules and configure the database
- **step 3** - populate the database with proteins
- **step 4** - start the WebPDB rest service
- **step 5** - start WebMOL and input the id of the desired protein

# Integration of PDB visualization service

