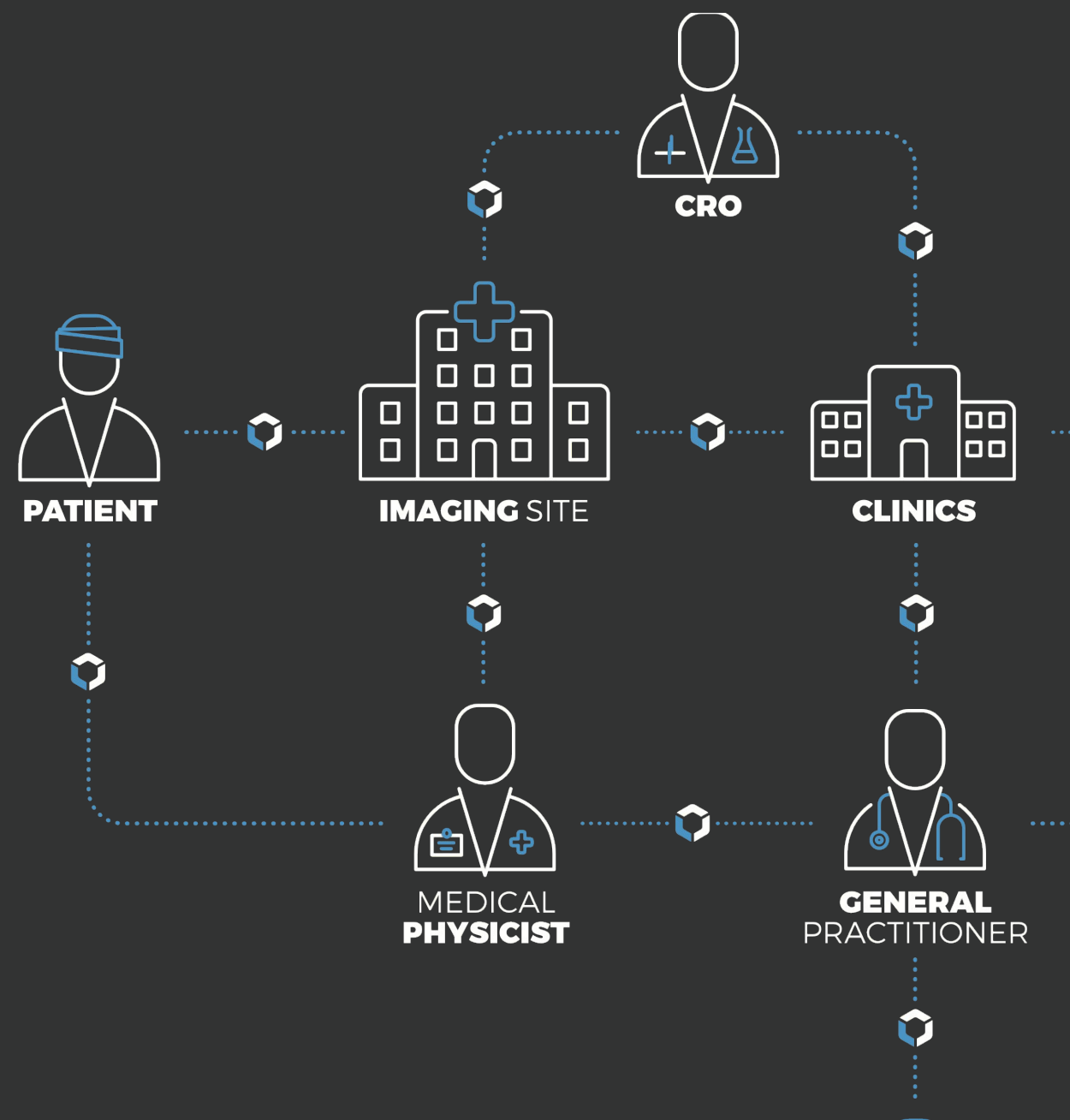


ORT_HANC

Using WebAssembly to render medical images

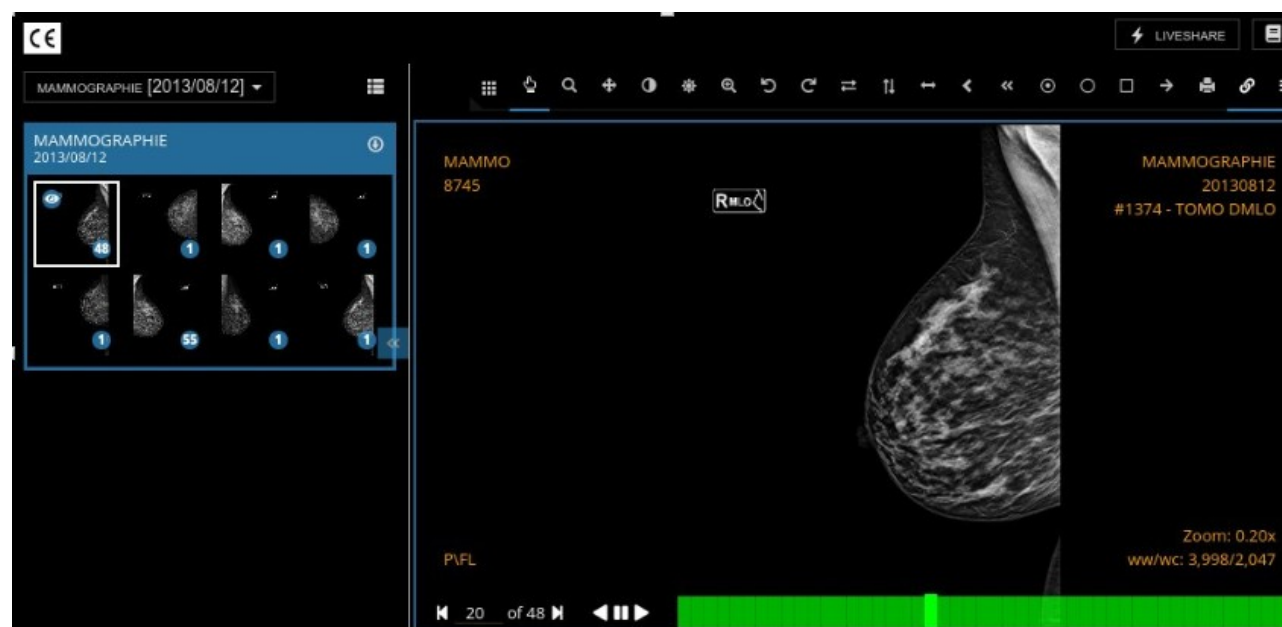


Free and open-source viewers for Orthanc

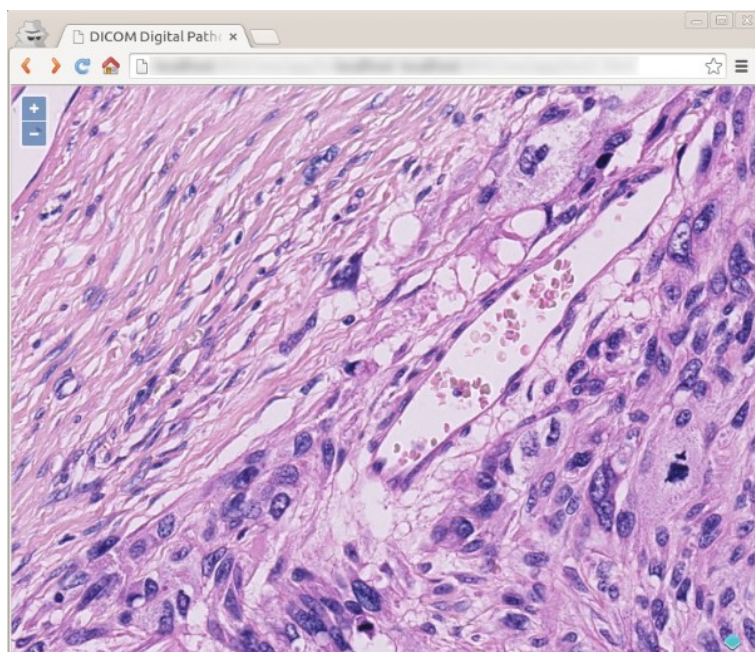
Basic reviewing



Advanced teleradiology (Osimis Web viewer)



Whole-slide imaging



External, Web:

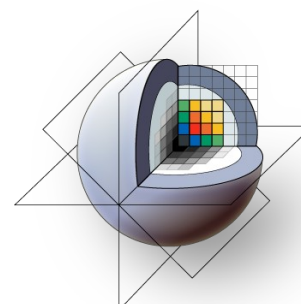
☐ ☐ Open Health
☐ ☐ Imaging Foundation

DWV, ...

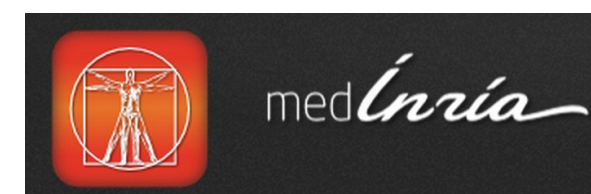
External, desktop:



Horos (Apple only)



3DSlicer



Aeksulap, ...

Two fully separate worlds



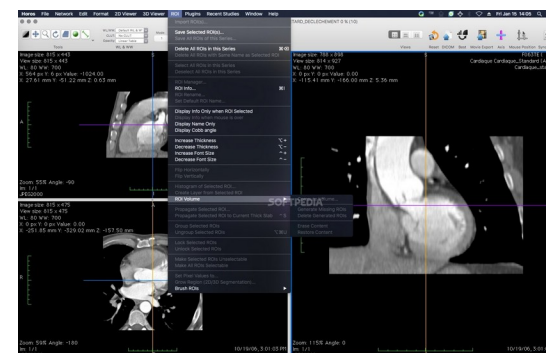
Web applications (teleradiology)

JavaScript + Cornerstone library



Desktop software (clinical radiology)

C++ +



The problems

No code reuse between Web and Desktop!
=> Fully redundant developments, separate teams

Desktop teams: How can I relocate some part of my software as a Web application for easy, fast delivery and to avoid the high cost of maintaining different ports and installers?

Web teams: How to use existing libraries for DICOM?

Question

Is it possible to run C++ client-side in Web browsers?



W3C[®]

Yes!



WEBASSEMBLY

moz://a



Google



redhat.

What is WebAssembly?

- **Bytecode for the Web**
- Open standard maintained by the W3C
- Official recommendation since 2019-12-05
- Precursors: Java applets, PNaCl from Google, asm.js from Mozilla...
- Supported by all the major Web browsers (including proprietary ones)



Official “C++ to WebAssembly” compiler

Hello, world! (1/2)

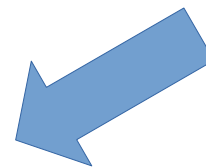


```
$ sudo apt install emscripten
```



```
#include <stdio.h>

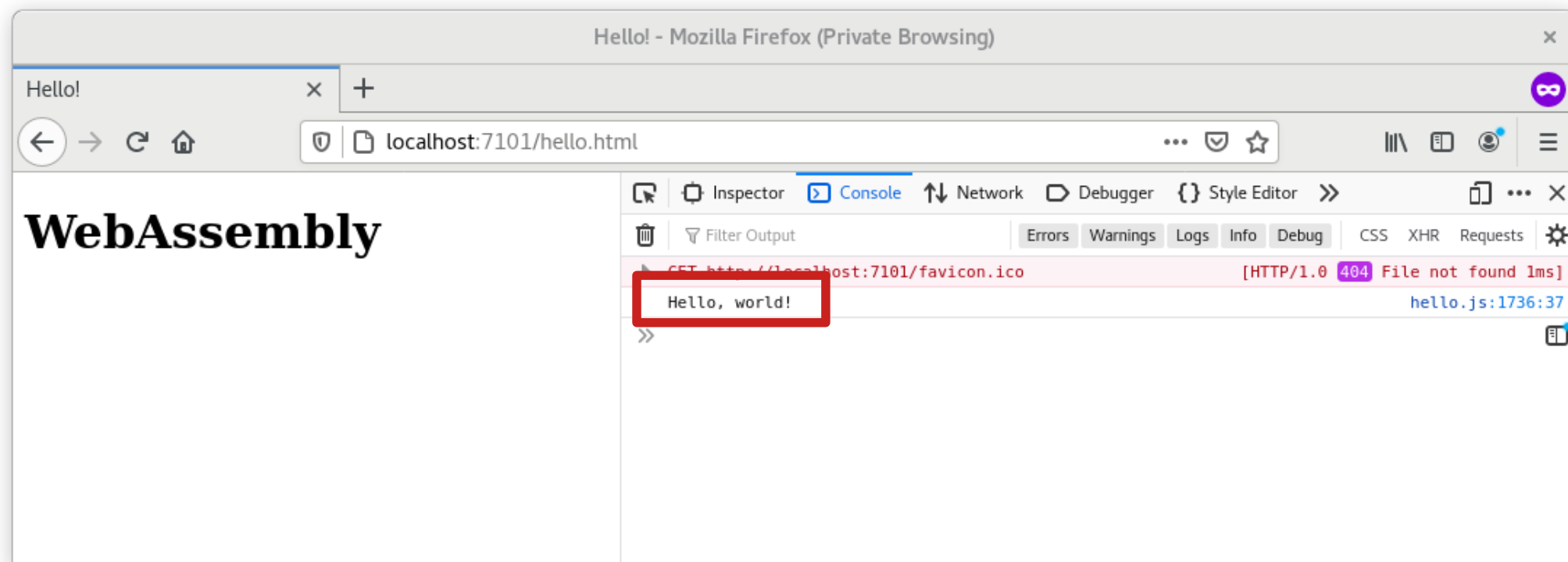
int main()
{
    printf("Hello, world!\n");
    return 0;
}
```



```
$ em++ ./hello.cpp -o hello.js
$ ls
hello.cpp    => C++ source code
hello.js     => JavaScript wrapper
hello.wasm   => WebAssembly bytecode
```


Hello, world! (2/2)

```
<!DOCTYPE html>
<html lang="en">
  <head>
    <meta charset="utf-8">
    <title>Hello!</title>
  </head>
  <body>
    <h1>WebAssembly</h1>
    <script src="hello.js" async></script>
  </body>
</html>
```



Stone of Orthanc

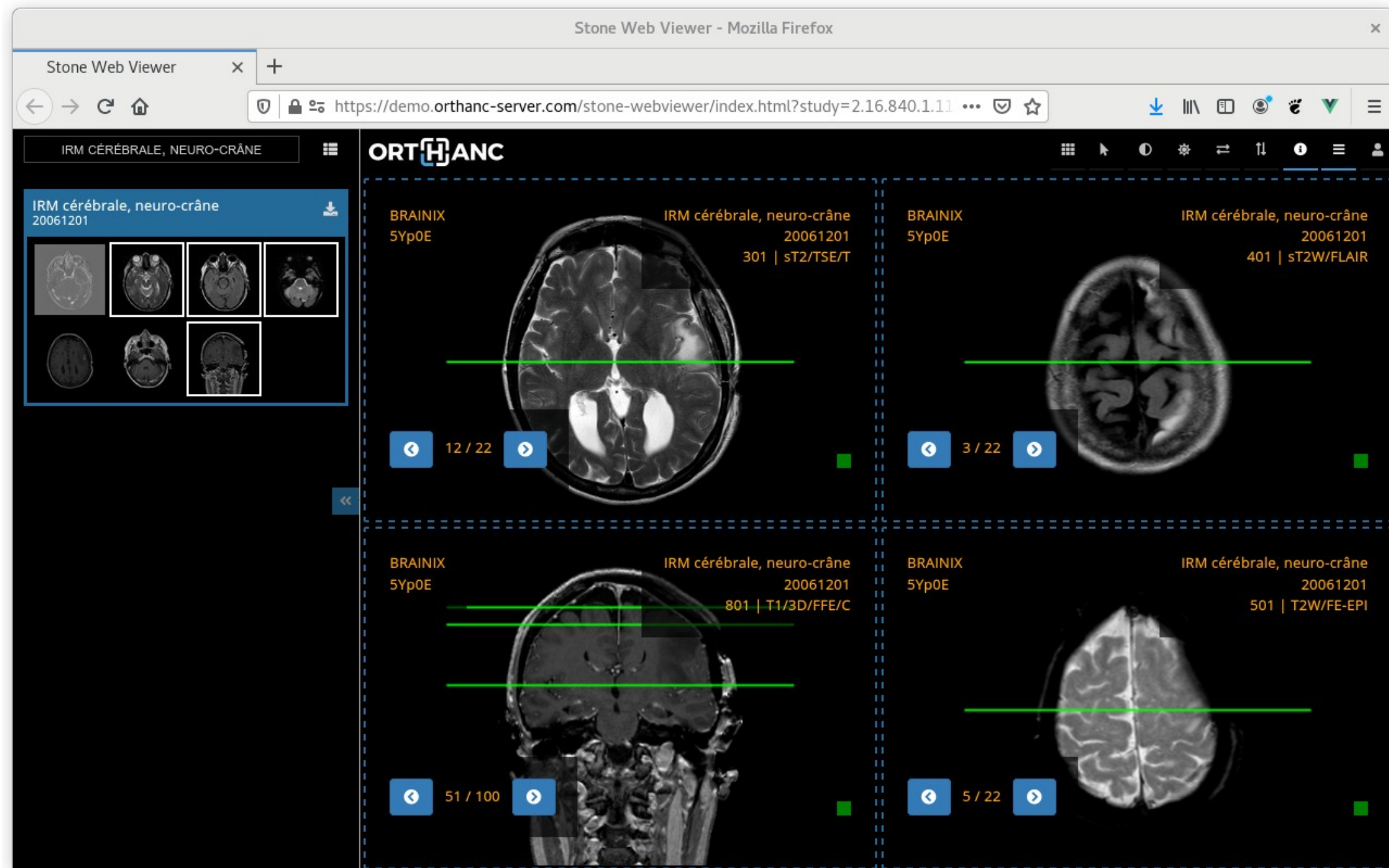
- **Lightweight, cross-platform C++ library to render medical images** (cf. VTK)
- Part of the Orthanc ecosystem
- Compatible with **WebAssembly**
- Compatible with **GUI libraries** (SDL, Qt...)
- Building block for the **Stone Web viewer**
- Obviously, libre software!

More features:

- *2D hardware acceleration (WebGL/OpenGL)*
- *Primitives for DICOM (parsing and DICOMweb)*
- *Built-in support of 3D volumes (MPR, volume reslicing)*
- *Support of oncology: PET-CT fusion, doses, contours...*

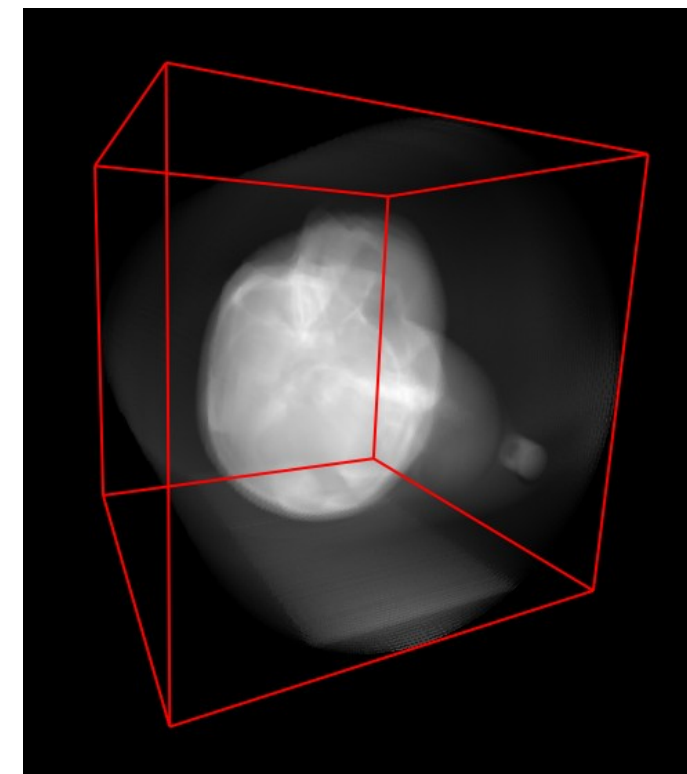
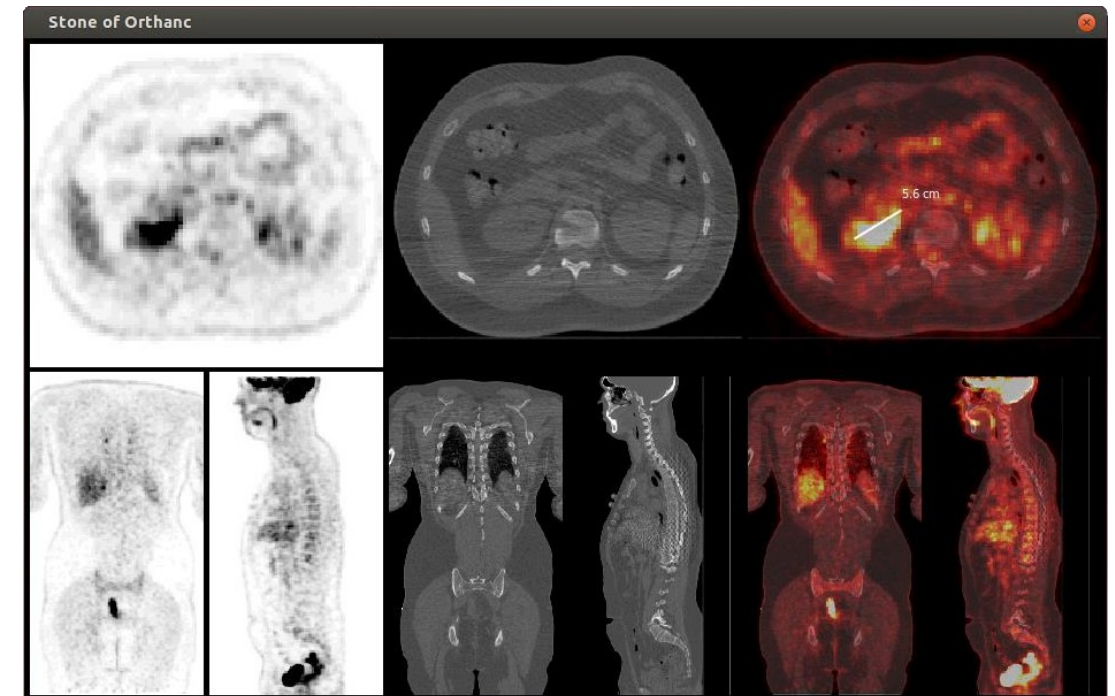
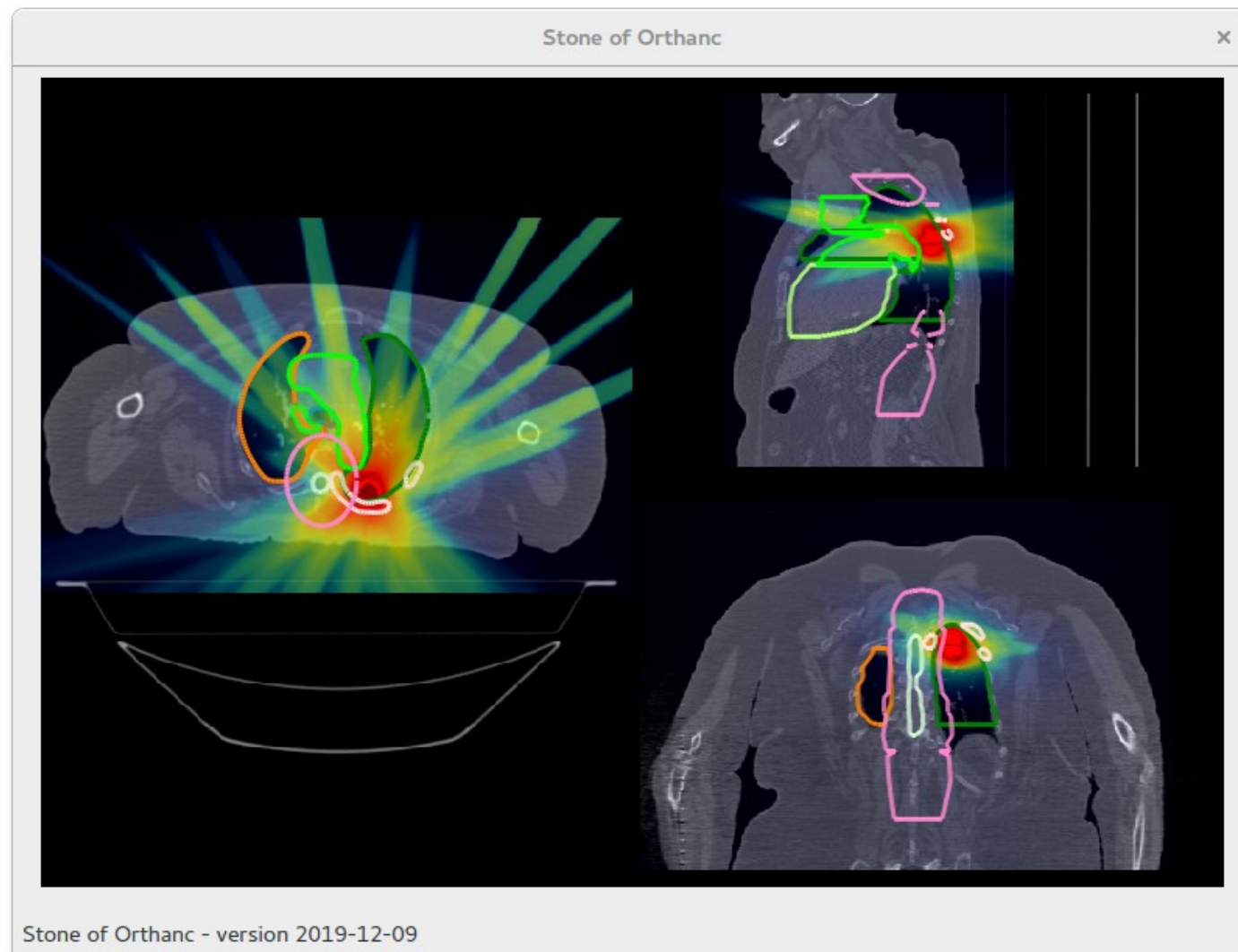


Stone Web viewer

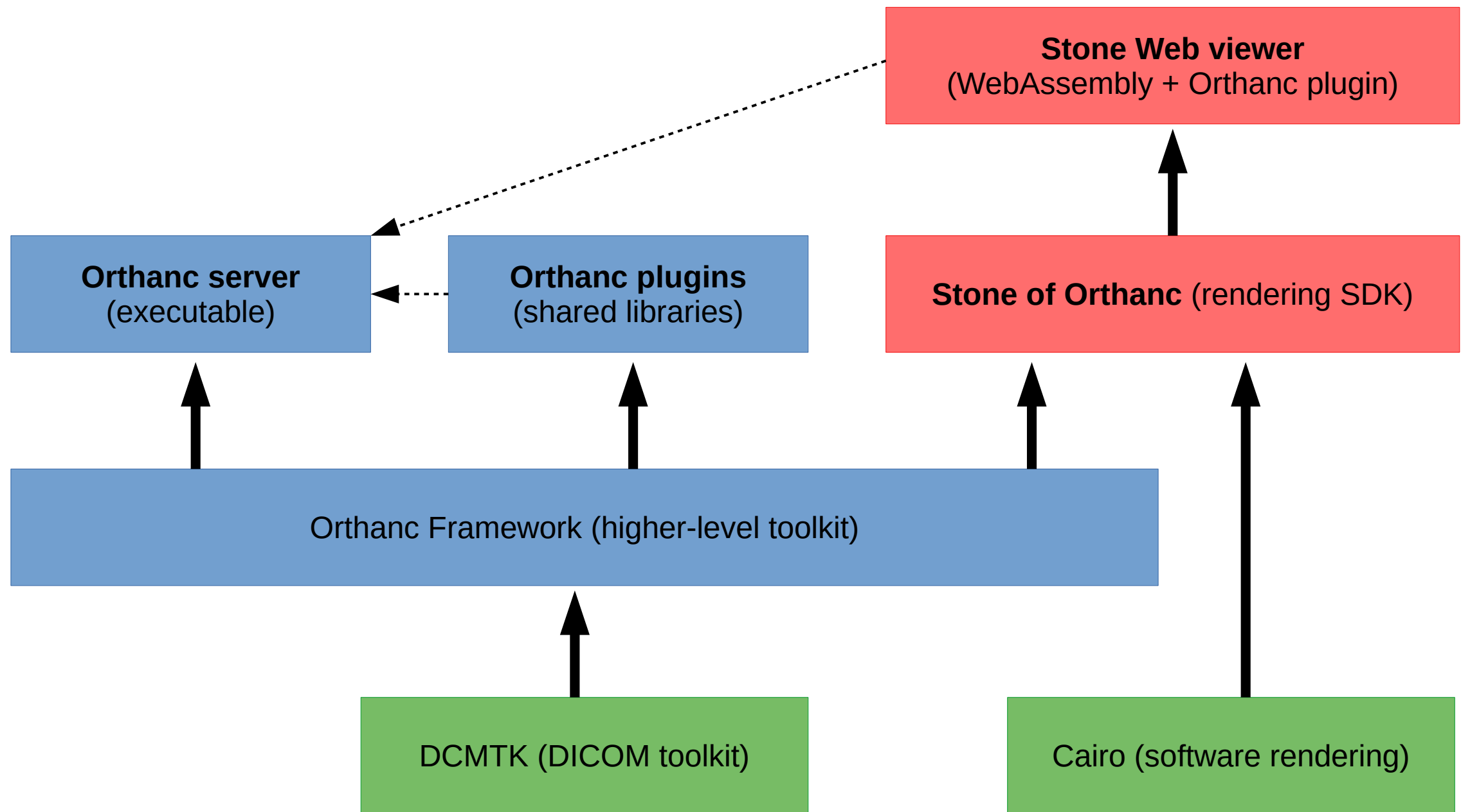


- Reuse the layout of the Osimis Web viewer (now deprecated)
- Online demo: <https://demo.orthanc-server.com/>
- Included in jodogne/orthanc-plugins and osimis/orthanc Docker images

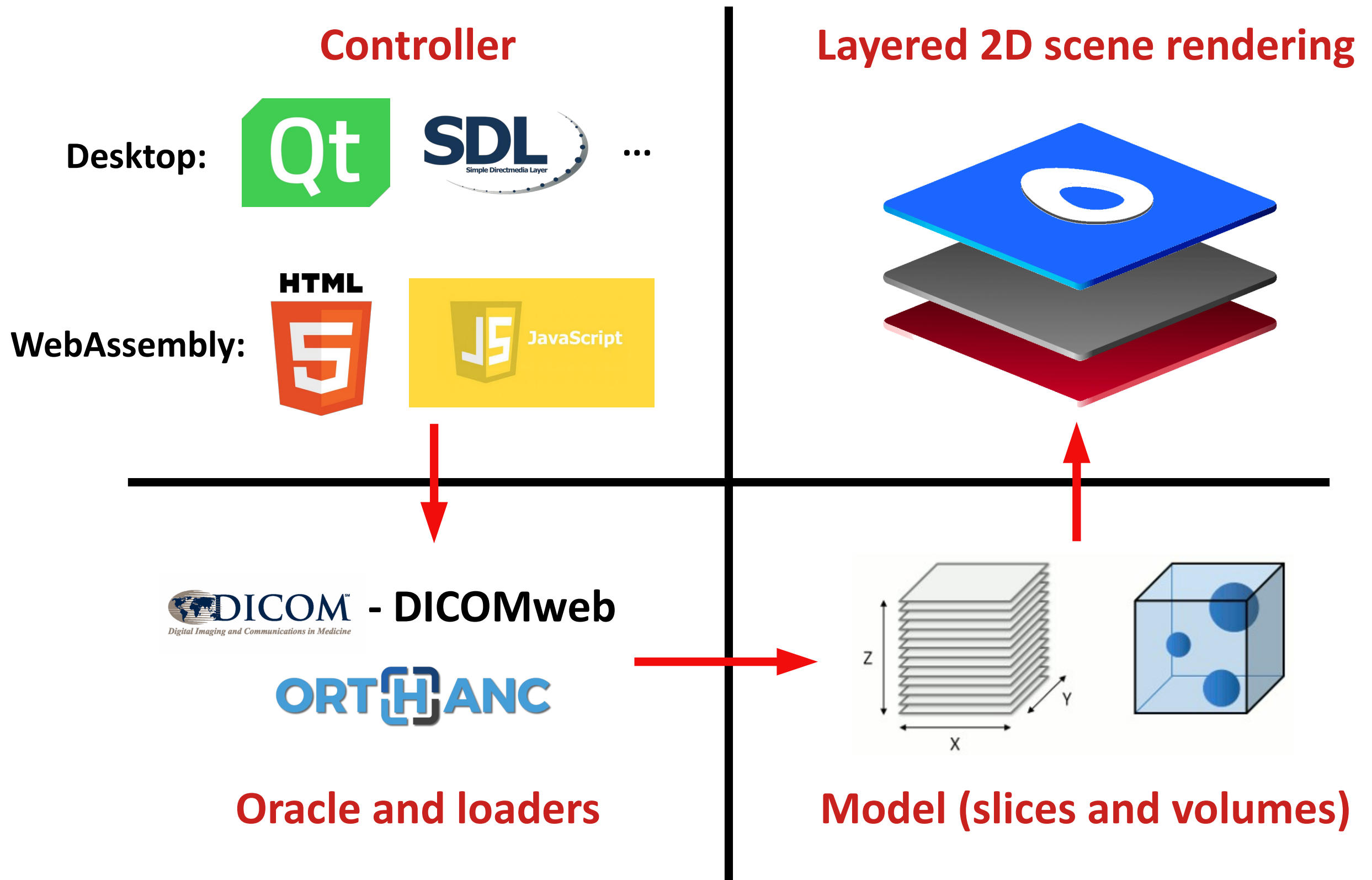
More advanced applications: 3D/MPR rendering



Place in the Orthanc ecosystem

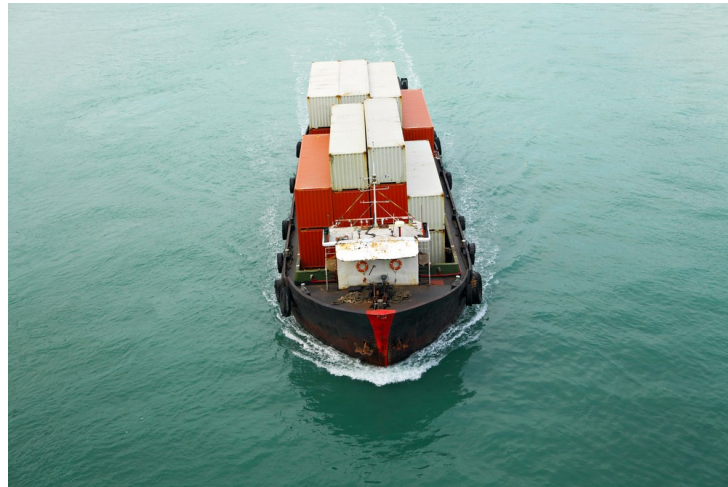


Overview of the Stone architecture



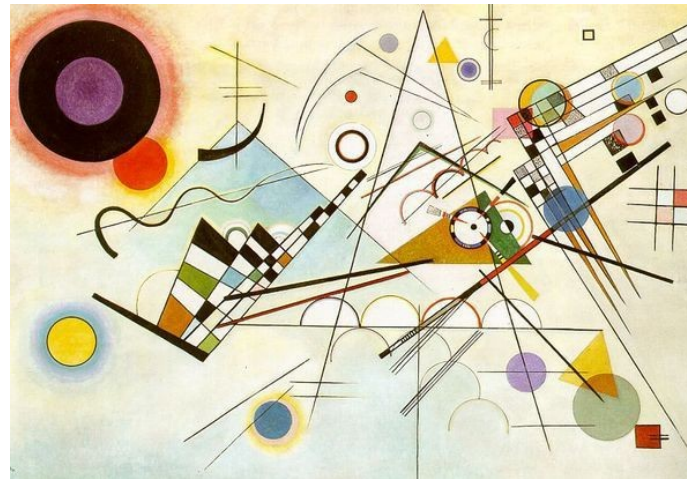
Challenges of Stone

Deployments



- Need a Web server to serve WebAssembly
- Complexity of interaction with many technologies (C++, HTML, JavaScript, DICOMweb...)
- Partial solution: **Orthanc plugins** can add routes in the HTTP server embedded into Orthanc

Different models



- JavaScript is single-threaded and promise-driven
- C++ is multi-threaded and sequential
- Management of windows differ strongly
- Solution: **Oracle that abstracts** the system and network primitives, plus **platform-specific 2D viewports**

Software libraries



- Need to compile each third-party library for WebAssembly (no repository of “side modules” so far)
- Few thought about packaging WebAssembly in GNU/Linux distros so far
- Solution: CMake scripts of Orthanc already knows how to **statically build** many libraries

Conclusions



Our mission statement:

“Freely share knowledge about medical imaging”

- The Orthanc ecosystem is also about displaying medical images!
- **Stone of Orthanc** is a lightweight, cross-platform C++ library
- **Stone Web viewer** combines Stone of Orthanc with WebAssembly
- The viewer can be used with other PACS servers than Orthanc (DICOMweb)
- First official release: December 2020!
- Easy integration with GNU Health: Simply open the URL of the study :-)