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ERROR

a pEdiatRic dosimetRy personalized platfQRm based on computational anthropomorphic phantoms

> Marie Skłodowska-Curie Actions Research and Innovation Staff RISE-Call: H2020-MSCA-RISE-2015 Project No.: 691203

Objectives

ERROR's objective is the development of a new software tool, which will offer to the clinician the possibility to assess imaging and therapeutic protocols predicting the absorbed dose per organ. This tool will be implemented and evaluated for pediatric applications, since this is a rather sensitive target group.



The Consortium

The **ERROR** project brings together a multidisciplinary consortium of specialists in different areas of medical physics, biomedical engineering, physicians, and computer engineers, who will join forces in order to design, implement, and clinically assess novel software tools. Two new SMEs will provide their expertise, and will exploit the project's outcome.

Project Partners	Role in Project
University of Patras	Project Coordination – Database –
(UPAT – GR)	MC Simulations – Clinical Assessment
Université de Bretagne Occidentale	MC Simulations – Clinical Assessment
(UBO – FR)	 – GPU Programming
Guy's and St Thomas NHS Foundation	Clinical Data – Clinical Assessment
Trust (GSTFT – GB)	Procedures
BET Solutions	Dissemination – MC Simulations –
(<i>BET – GR</i>)	Software Development
LIBRA MLI Ltd.	Machine Learning – Software
(LIBRA – GB)	Development

Methodology & Workflow

Based on clinical data, and on the available literature we aim to quantify personalized dosimetry on the most critical organs in children (2-15 years old), and create a universal tool for the total calculation of the absorbed dose per organ, according to the:

- Specified radiopharmaceutical biodistributions
- Injected activity
- X-ray model irradiation
- Brachytherapy source model and irradiation type
- Anatomical characteristics (age, weight, height, gender)

The creation of the dosimetric software will be based on:

- Accurate physical processes modeling
- Realistic MC simulations incorporating biokinetics
- High resolution anthropomorphic computational models
- Powerful computational resources (grids, GPU, HPC)

7 Working Packages are connected towards this effort.

