


Introducción al desarrollo de aplicaciones de IA

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Quibim SL

INTELIGENCIA ARTIFICIAL

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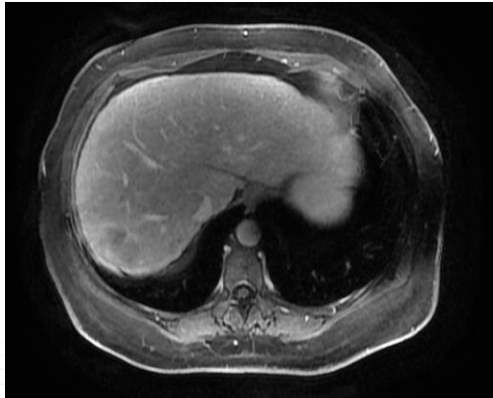
- Creación de la base de datos
 - Anotación
 - Estrategias de entrenamiento:
 - Modelos basados en características
 - Modelos basados en imagen
 - Estrategias de validación
 - Despliegue de modelos de IA
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 - Anotación
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 - Estrategias de validación
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- 

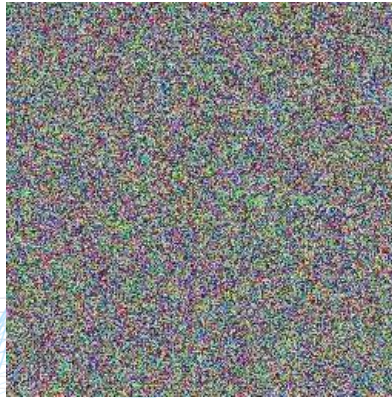
Creación de la base de datos

Imagen original



Tumor

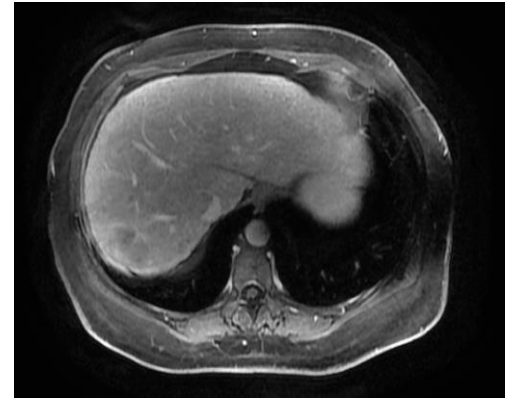
Ruido



+

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Imagen con ruido

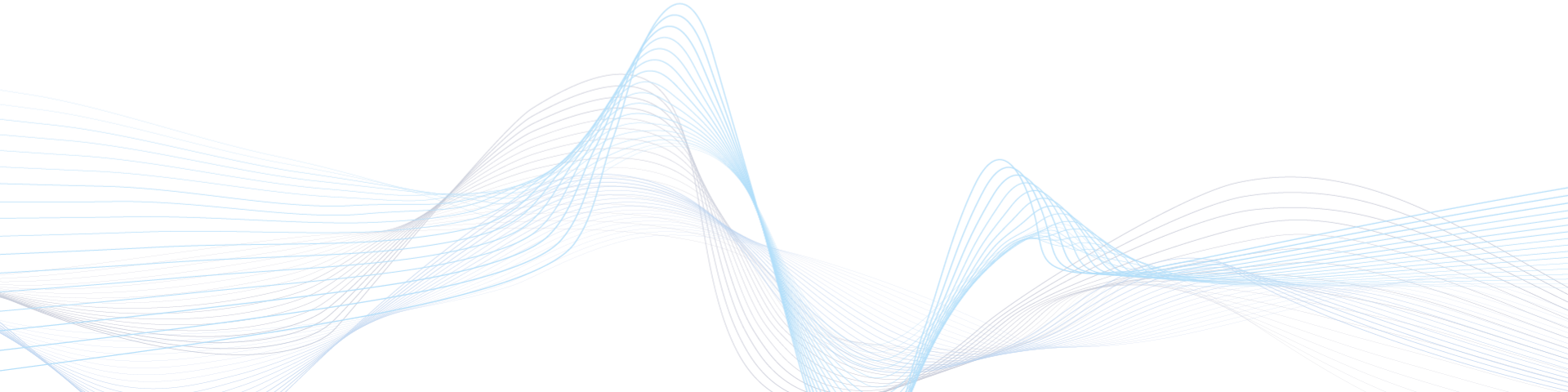


Sano

Los modelos de IA son muy sensibles a pequeñas variaciones de la imagen de entrada

Creación de la base de datos


Requerimientos para el desarrollo de un modelo de IA robusto:



Creación de la base de datos

Requerimientos para el desarrollo de un modelo de IA robusto:

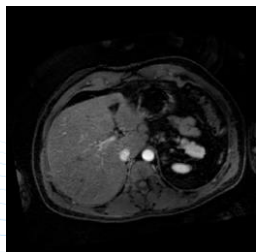
1. Base de datos heterogénea:

- Mujeres y hombres
 - Diferentes edades
 - Diferentes nacionalidades
 - Multi-céntrico
 - Diferentes escáneres
 - Diferentes protocolos de adquisición
 - ...
- 

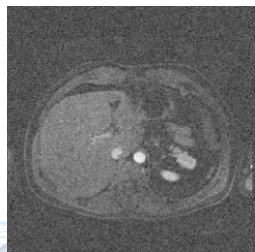
Creación de la base de datos

Requerimientos para el desarrollo de un modelo de IA robusto:

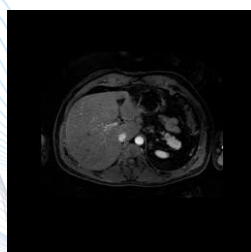
1. Base de datos heterogénea
2. **Técnicas de aumento de datos:**



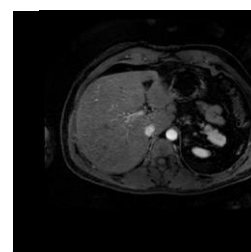
Rotaciones



Ruido



Zoom



Translaciones

...

Creación de la base de datos

Requerimientos para el desarrollo de un modelo de IA robusto:

1. Base de datos heterogénea
2. Técnicas de aumento de datos
3. **Validación externa**

Design Characteristics of Studies Reporting the Performance of Artificial Intelligence Algorithms for Diagnostic Analysis of Medical Images: Results from Recently Published Papers

Dong Wook Kim, MD^{1*}, Hye Young Jang, MD^{2*}, Kyung Won Kim, MD, PhD², Youngbin Shin, MS², Seong Ho Park, MD, PhD²

¹Department of Radiology, Taean-gun Health Center and County Hospital, Taean-gun, Korea; ²Department of Radiology and Research Institute of Radiology, University of Ulsan College of Medicine, Asan Medical Center, Seoul, Korea


Results: Of 516 eligible published studies, only 6% (31 studies) performed external validation. None of the 31 studies adopted all three design features: diagnostic cohort design, the inclusion of multiple institutions, and prospective data collection for external validation. No significant difference was found between medical and non-medical journals.

Creación de la base de datos

Requerimientos para el desarrollo de un modelo de IA robusto:

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Checklist for Artificial Intelligence in Medical Imaging (CLAIM): A Guide for Authors and Reviewers

 John Mongan,  Linda Moy,  Charles E. Kahn, Jr 

✓ **Author Affiliations**

Published Online: Mar 25 2020 | <https://doi.org/10.1148/ryai.2020200029>

Checklist for Artificial Intelligence in Medical Imaging (CLAIM)		
Section/Topic	Nº.	Item
TITLE or ABSTRACT	1	Identification as a study of AI methodology, specifying the category of technology used (eg, deep learning)
	2	Structured summary of study design, methods, results, and conclusions
ABSTRACT	3	Scientific and clinical background, including the intended use and clinical role of the AI approach
	4	Study objectives and hypotheses
INTRODUCTION	5	Prospective or retrospective study
	6	Study goal, such as model creation, exploratory study, feasibility study, noninferiority trial
METHODS	7	Data sources
	8	Eligibility criteria: how, where, and when potentially eligible participants or studies were identified (eg, symptoms, results from previous tests, inclusion in registry, patient-care setting, location, dates)
Study Design	9	Data preprocessing steps
	10	Selection of data subsets, if applicable
Data	11	Definitions of data elements, with references to common data elements
	12	De-identification methods
Ground Truth	13	How missing data were handled
	14	Definition of ground truth reference standard, in sufficient detail to allow replication
Model	15	Rationale for choosing the reference standard (if alternatives exist)
	16	Source of ground truth annotations; qualifications and preparation of annotators
Training	17	Annotation tools
	18	Measurement of inter- and intratester variability; methods to mitigate variability and/or resolve discrepancies
Evaluation	19	Intended sample size and how it was determined
	20	How data were assigned to partitions; specify proportions
RESULTS	21	Level at which partitions are disjoint (eg, image, study, patient, institution)
	22	Detailed description of model, including inputs, outputs, all intermediate layers and connections
DISCUSSION	23	Software libraries, frameworks, and packages
	24	Initialization of model parameters (eg, randomization, transfer learning)
OTHER INFORMATION	25	Details of training approach, including data augmentation, hyperparameters, number of models trained
	26	Method of selecting the final model
DISCUSSION	27	Ensembling techniques, if applicable
	28	Metrics of model performance
DISCUSSION	29	Statistical measures of significance and uncertainty (eg, confidence intervals)
	30	Robustness or sensitivity analysis
DISCUSSION	31	Methods for explainability or interpretability (eg, saliency maps) and how they were validated
	32	Validation or testing on external data
DISCUSSION	33	Flow of participants or cases, using a diagram to indicate inclusion and exclusion
	34	Demographic and clinical characteristics of cases in each partition
DISCUSSION	35	Performance metrics for optimal model(s) on all data partitions
	36	Estimates of diagnostic accuracy and their precision (such as 95% confidence intervals)
DISCUSSION	37	Failure analysis of incorrectly classified cases
	38	Study limitations, including potential bias, statistical uncertainty, and generalizability
DISCUSSION	39	Implications for practice, including the intended use and/or clinical role
	40	Registration number and name of registry
DISCUSSION	41	Where the full study protocol can be accessed
	42	Sources of funding and other support; role of funders

Creación de la base de datos

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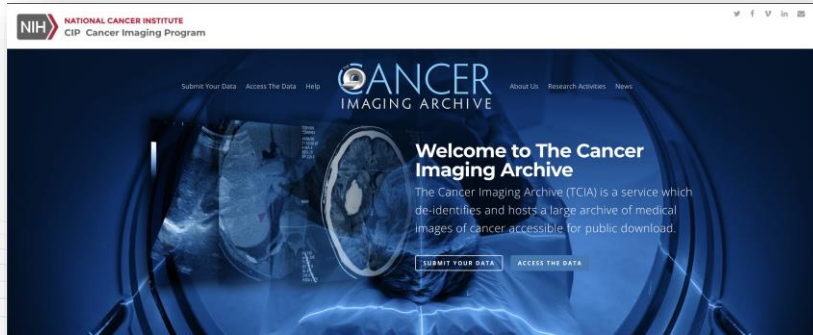
Certificados para poder vender como producto sanitario



Creación de la base de datos

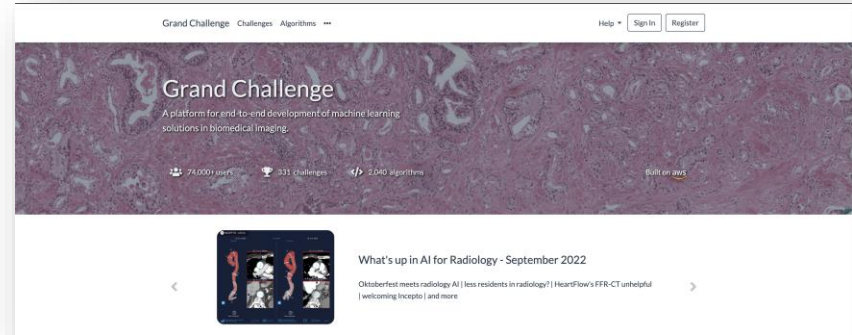
Bases de datos públicas

The Cancer Imaging Archive (TCIA)




<https://www.cancerimagingarchive.net/>

Grand Challenge



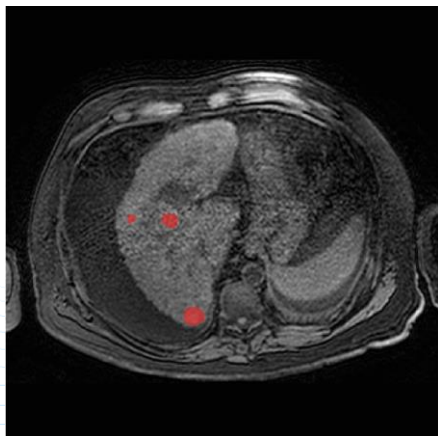
<https://grand-challenge.org/>

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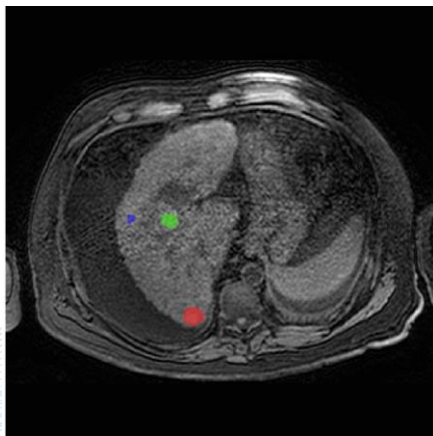
- Creación de la base de datos
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Anotación

Segmentación

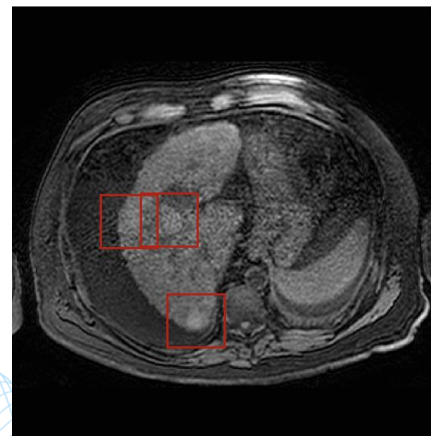


Segmentación
semántica



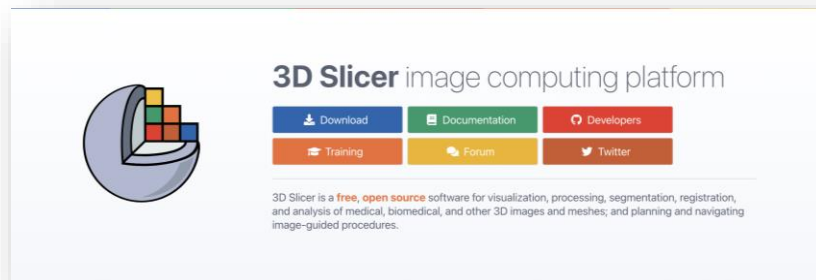
Segmentación de
instancias

Detección de objetos / Localización



Anotación

Herramientas



<https://www.slicer.org/>



<http://www.itksnap.org/pmwiki/pmwiki.php>

Anotación

Variabilidad intra- inter- observador → Guías de anotación y consenso

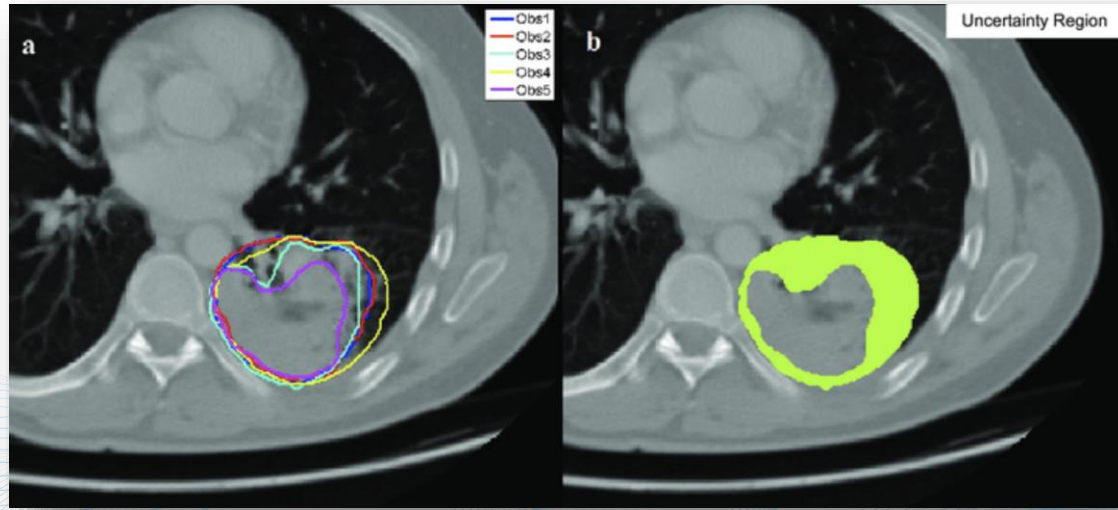


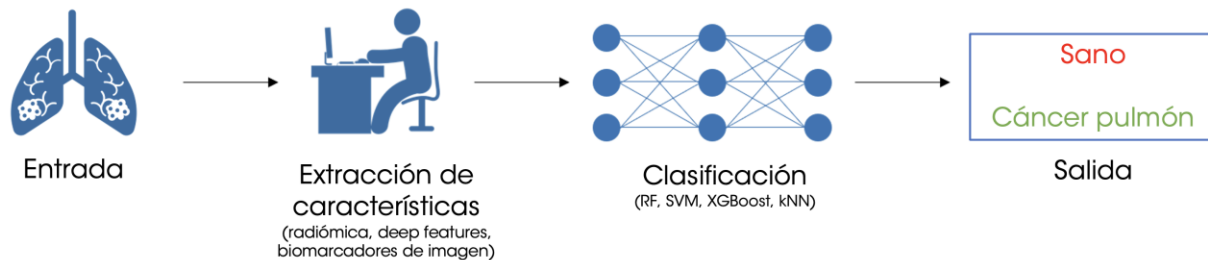
Image source: Rios Velazquez et al. Volumetric CT-based segmentation of NSCLC using 3D-slicer.

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Estrategias de entrenamiento

Modelos basados en características



Modelos basados en imagen



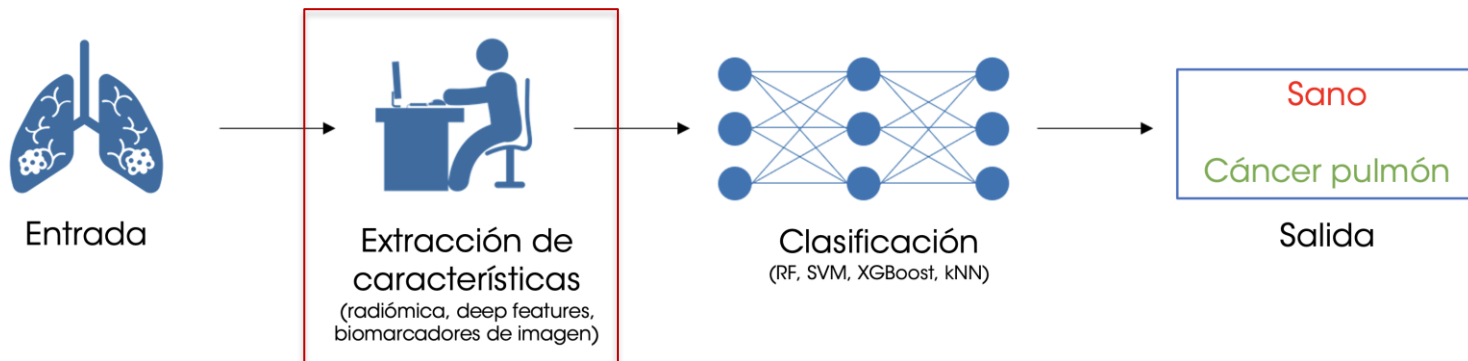
Datos de entrenamiento

-

+

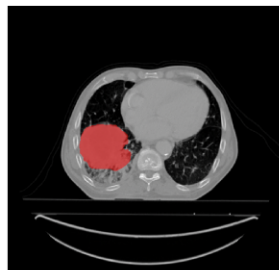
Estrategias de entrenamiento

Modelos basados en características



Estrategias de entrenamiento

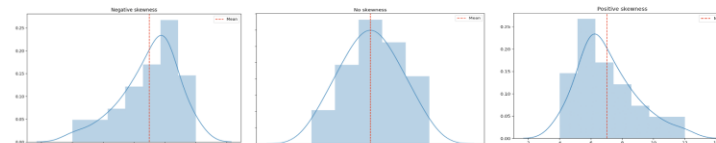
Modelos basados en características *Características de radiómica*



**Características para el estudio de la
heterogeneidad del tumor.**

PyRadiomics*

1. First order: Distribution of individual voxel values



2. Second order: Statistical inter-relationships between
neighboring voxels

GLCM GLRLM GLSZM GLDM NGTDM

3. Shape: Geometric properties of the delineated ROI



4. Higher order features

Square, exponential, logarithm, wavelet, LoG

*van Griethuysen, J. J. M. et al. (2017). Computational Radiomics System to Decode the Radiographic Phenotype. Cancer Research

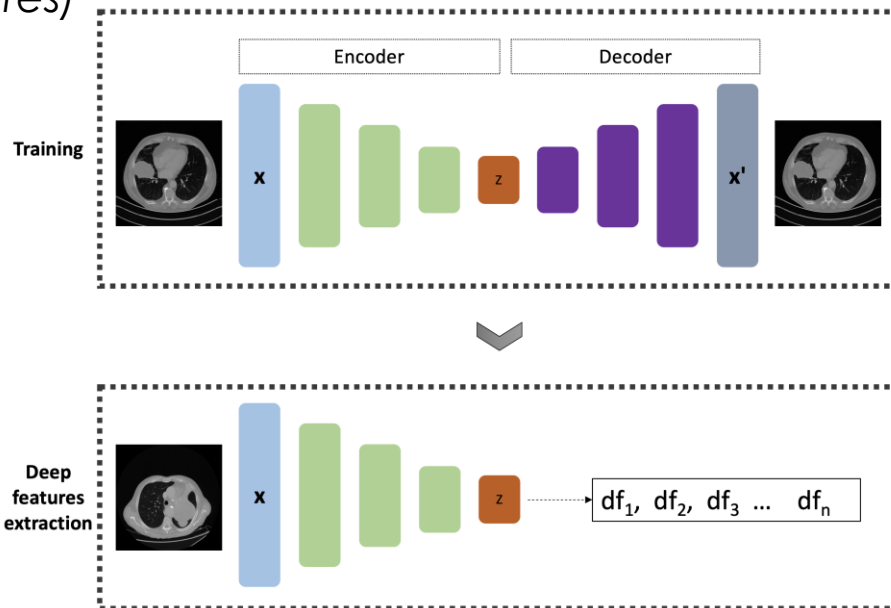
Estrategias de entrenamiento

Modelos basados en características

Características profundas (Deep features)

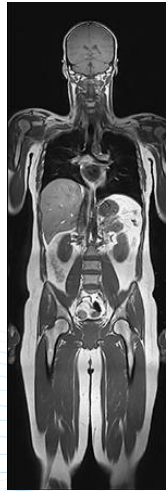
Se emplea la capacidad de aprendizaje de las **CNN**. Así, se entrena una **autocodificador** capaz de extraer la **información más relevante** de la imagen de entrada (características).

Una vez entrenado, se emplea el **codificador** para extraer dichas **características** de imágenes nuevas.



Estrategias de entrenamiento

Modelos basados en características *Biomarcadores de imagen*



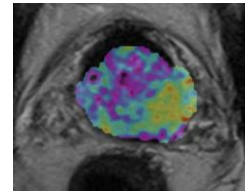
RM



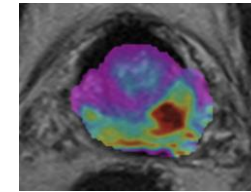
TC



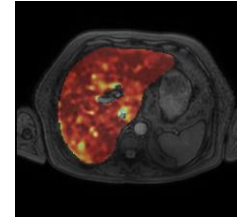
PET



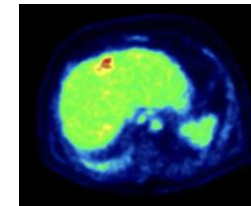
ADC



K^{trans}



PDFF



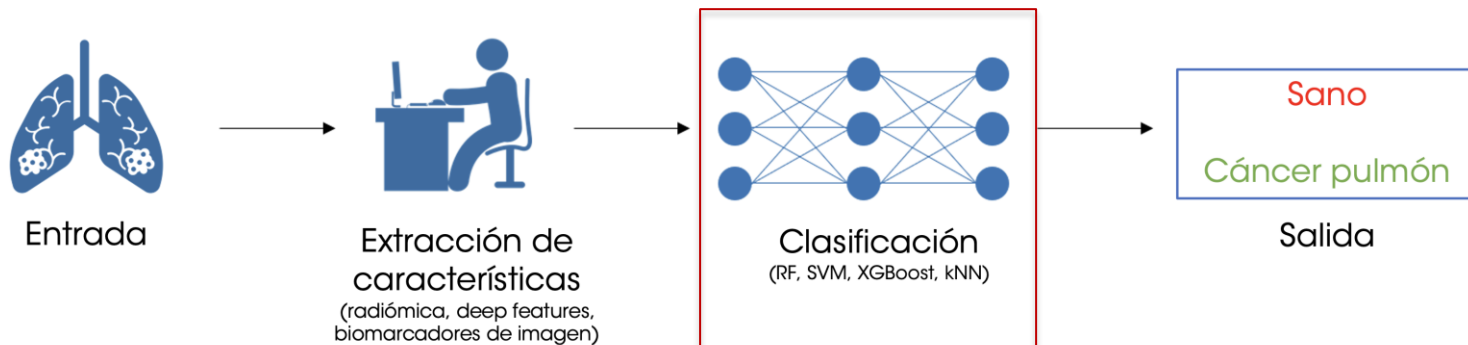
SUV

...

Fuente imágenes: ADC, K^{trans} and PDFF (Quibim SL, Valencia, Spain). SUV (Dimitrakopoulou-Strauss, A. et al. Kinetic modeling and parametric imaging with dynamic PET for oncological applications: general considerations, current clinical applications, and future perspectives. Eur J Nucl Med Mol Imaging (2021)

Estrategias de entrenamiento

Modelos basados en características



Estrategias de entrenamiento

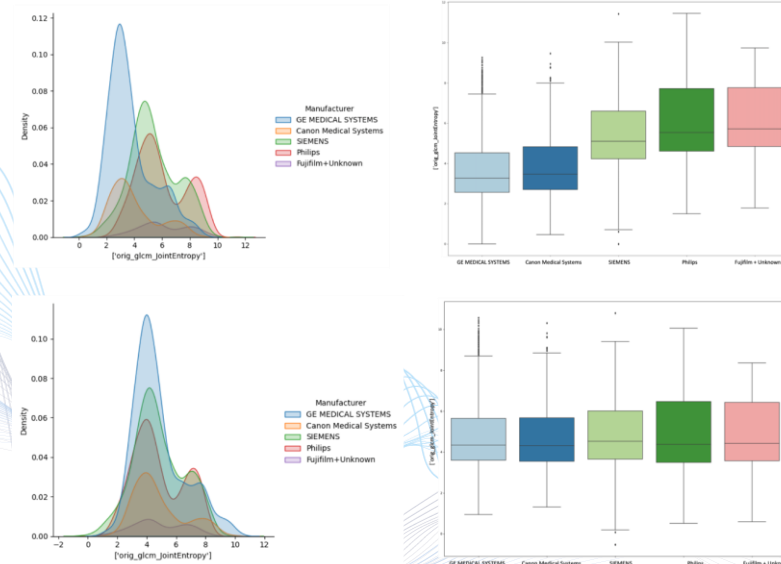
Modelos basados en características

Entrenamiento del modelo

1. Armonización de características

Las técnicas de armonización permiten reducir las diferencias entre fabricantes y protocolos de adquisición.

ComBat* es una técnica extendida que se empleó previamente en genómica.

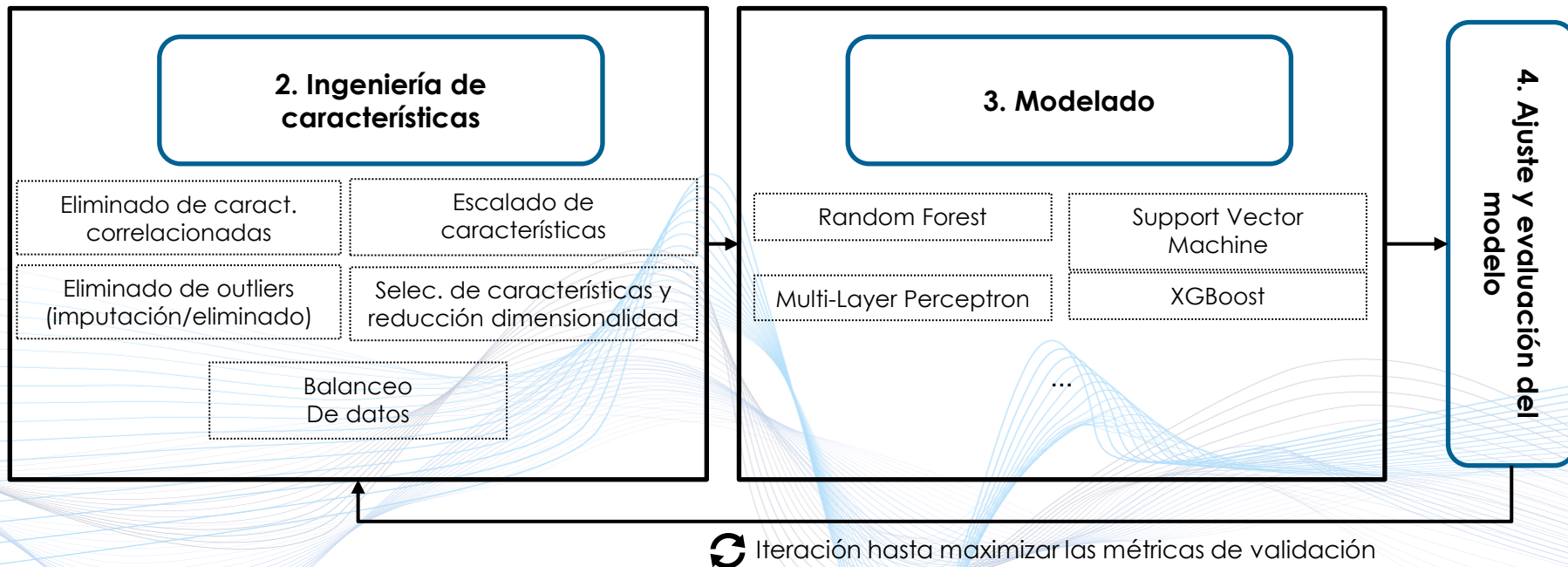


*Johnson WE et. al. Adjusting batch effects in microarray expression data using empirical Bayes methods. *Biostatistics*. (2007)

Estrategias de entrenamiento

Modelos basados en características

Entrenamiento del modelo



Estrategias de entrenamiento

Modelos basados en imagen



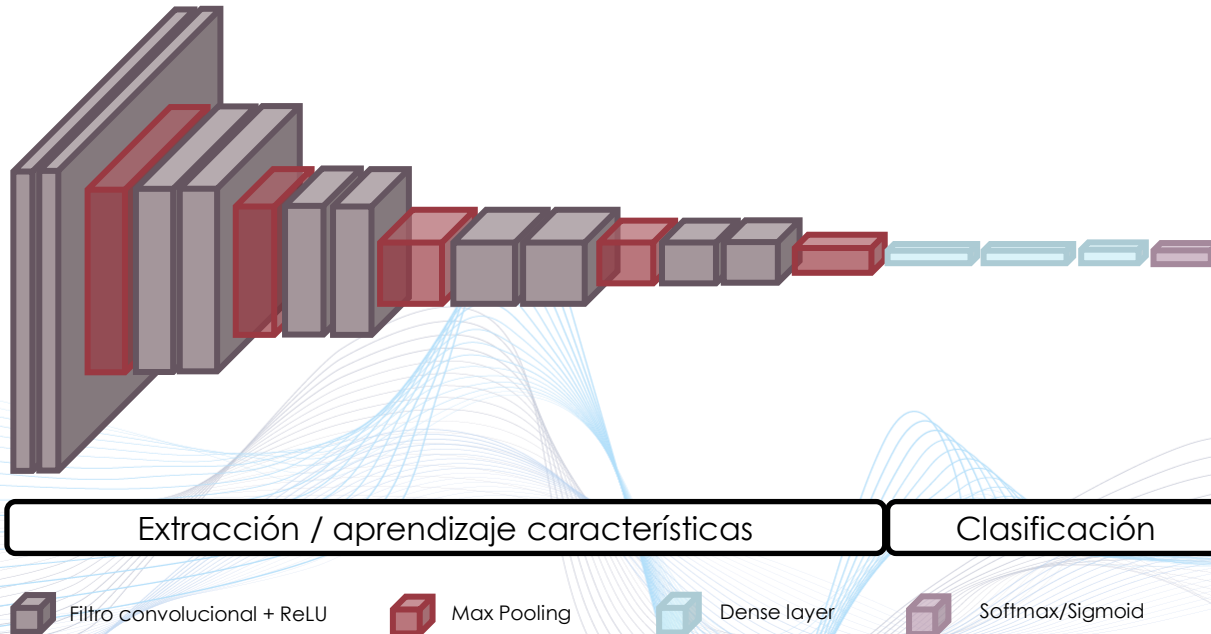
Entrada



Salida

Estrategias de entrenamiento

Modelos basados en imagen



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Estrategias de validación

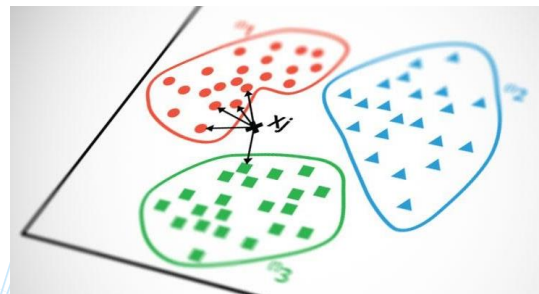
Validación técnica

Comparación de los resultados contra una referencia (fantoma, DRO, dispositivo de referencia, anotaciones, etc.). Se emplea para la validación de los algoritmos de extracción de características.



Validación clínica

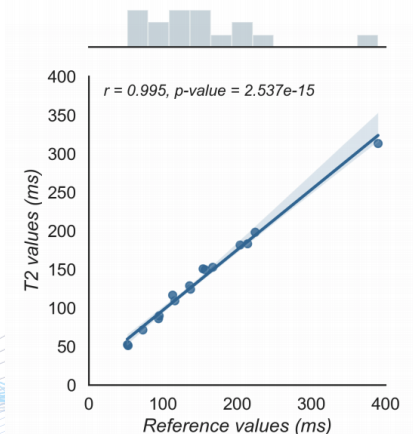
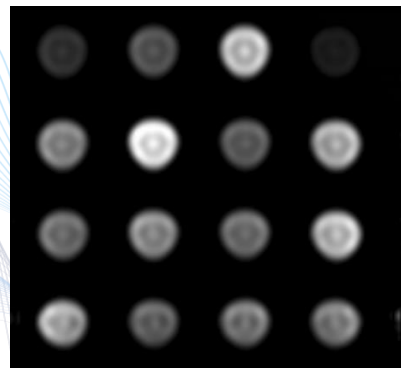
Correlación con un objetivo clínico (e.g., estadiaje, respuesta al tratamiento, tiempo de supervivencia, etc.).



Estrategias de validación

Validación técnica. Fantoma

Objeto físico compuesto por tubos con valores conocidos (e.g., T1 y T2 en RM).

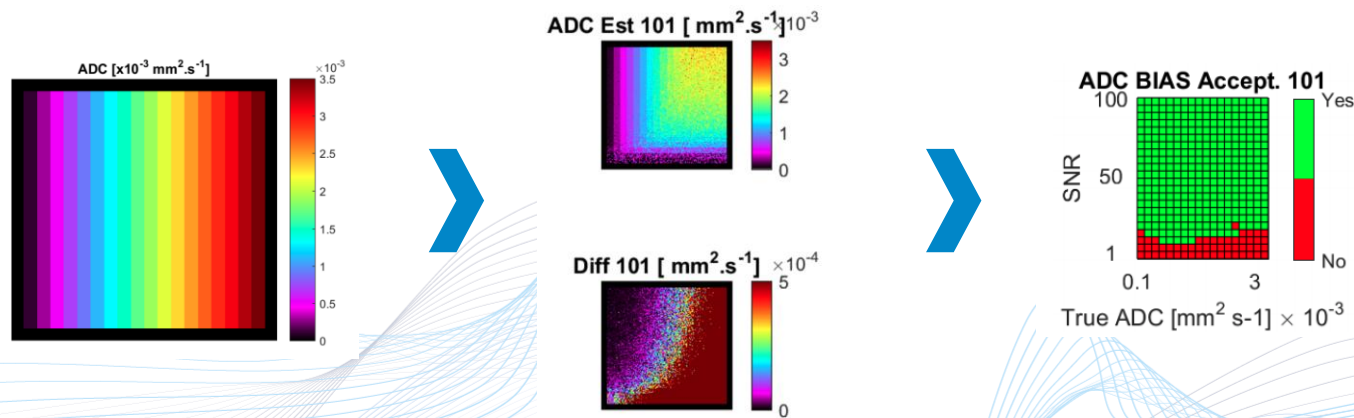


Validación de un algoritmo para la cuantificación del T2.

Estrategias de validación

Validación técnica. Objeto de referencia digital (DRO)

Imagen con los valores cuantitativos de referencia

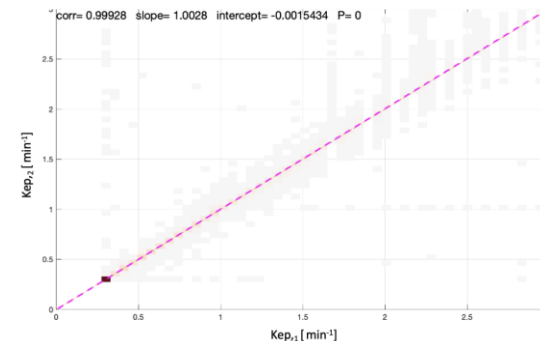
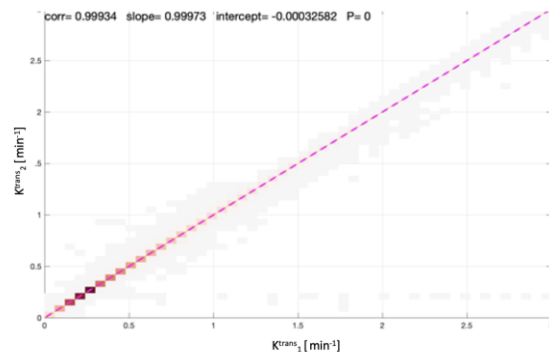
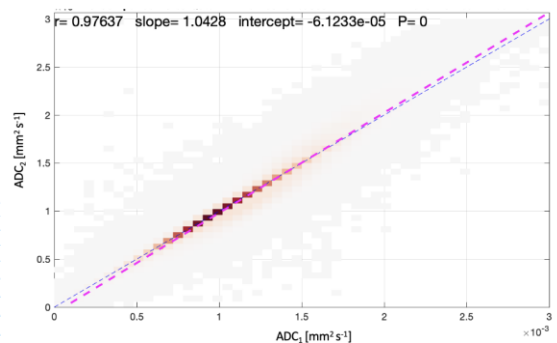


Perfil de QIBA : Imagen potenciada en diffusion (DWI). Validación de un algoritmo de cuantificación del coeficiente aparente de diffusion (ADC)

Estrategias de validación

Validación técnica. Dispositivo de referencia

Comparativa con un dispositivo médico previamente validado.

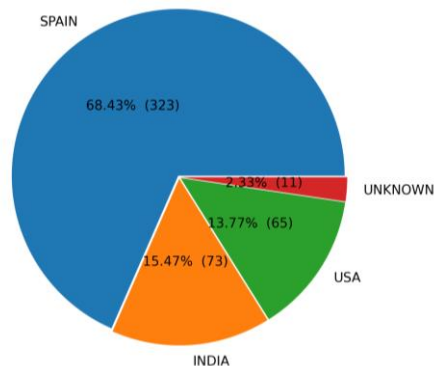


Estrategias de validación

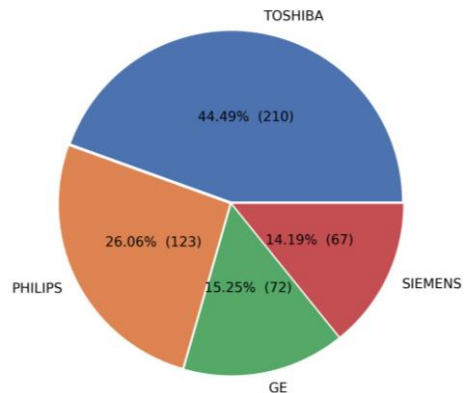
Validación técnica. Anotaciones expertos

Comparativa con datos anotados manualmente y revisados por expertos.
Para garantizar la robustez del modelo hay que evaluarlo en diferentes escenarios (origen, fabricantes, campo magnético, etc.)

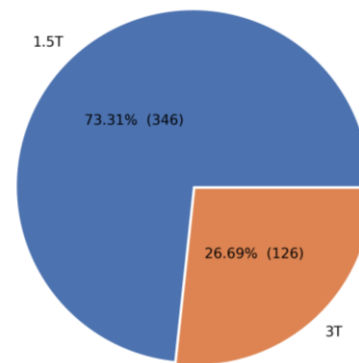
ORIGIN OF THE DATA (N = 472)



IMAGING VENDORS (N = 472)

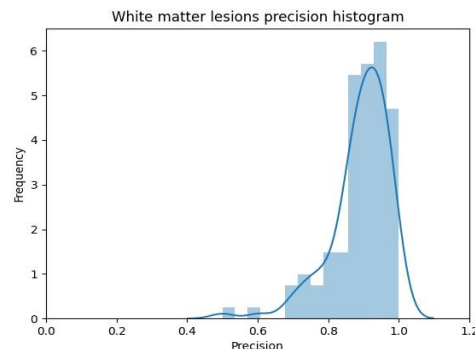
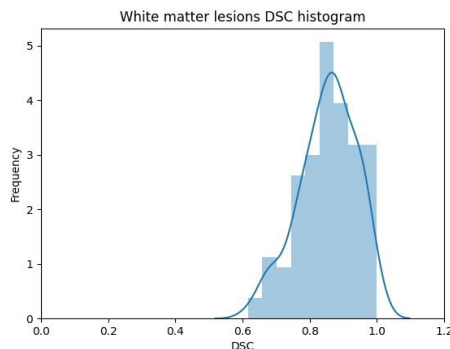


MAGNETIC FIELD (N = 472)



Estrategias de validación

Validación técnica. Anotaciones expertos. Segmentación



Métricas	Media	Std	Mediana
DSC	0.85 (0.84–0.87)	0.09 (0.06–0.10)	0.87 (0.85–0.88)
Precision	0.89 (0.88–0.91)	0.08 (0.07–0.10)	0.91 (89–0.92)
Recall	0.81 (0.79–0.83)	0.12 (0.10–0.14)	0.82 (0.80–0.84)

Validación de un algoritmo de segmentación automática de hiperintensidades de sustancia blanca.

Estrategias de validación

Validación clínica

Comparación contra un objetivo clínico:

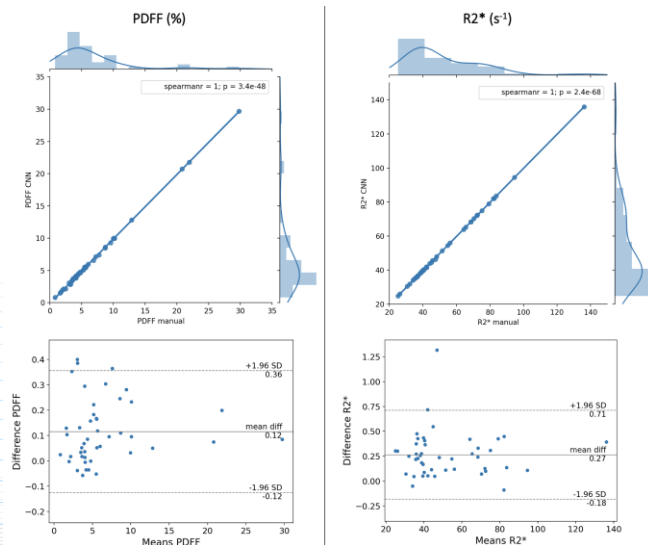
- **Modelos diagnósticos:** comparativa frente a estándar de referencia (e.g., biopsia). Estrategias de superioridad o no inferioridad.
- **Modelos predictivos:** análisis de supervivencia (e.g., supervivencia, tiempo libre de progresión), modelos de clasificación (e.g., response al tratamiento).



Estrategias de validación

Validación clínica

Valores cuantitativos: Segmentación CNN vs manual



Valores de PDF vs. Grados esteatosis (biopsia)

Segmentation	Grades	Cut-off	AUC	95% CI	Sn	Sp
PDF – Steatosis (%)						
ROI	S _{≥1}	> 7.7	0.96	0.93-0.98	94	84
VOI		> 7.8	0.97	0.94-0.99	95	84
ROI	S _{≥2}	> 10.2	0.95	0.92-0.98	92	90
VOI		> 10.8	0.96	0.93-0.99	88	91
ROI	S _{≥3}	> 12.8	0.96	0.93-0.99	93	87
VOI		> 14.1	0.96	0.93-0.99	89	91

Validación de un algoritmo para la cuantificación de PDF y R2* hepáticos

Estrategias de validación

Validación clínica

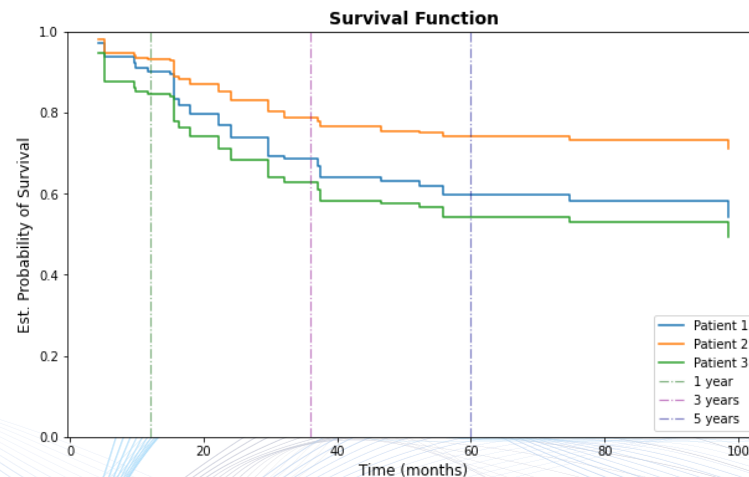
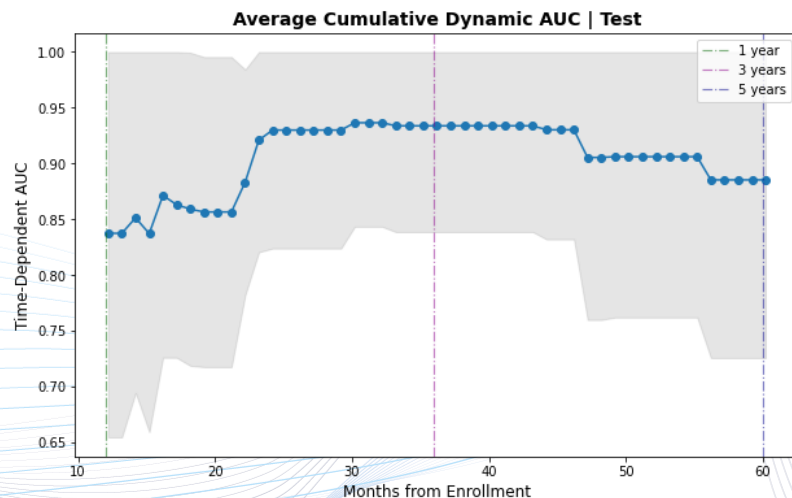
Validación de un algoritmo para la predicción del tiempo de supervivencia en pacientes con neuroblastoma

Model	Modifications	C. Index IPCW		C. Index		Mean AUC		Mean Brier Score	
		Train	Test	Train	Test	Train	Test	Train	Test
Cox	Boruta	0.745	0.709	0.785	0.728	0.829	0.875	0.108	0.117
ElasticNet Cox	Boruta	0.745	0.709	0.785	0.729	0.830	0.875	0.108	0.117
Random Survival Forest	PCA	0.913	0.721	0.928	0.743	0.962	0.895	0.085	0.123
Extra Survival Trees	PCA	0.869	0.718	0.893	0.747	0.928	0.827	0.093	0.127
Gradient Boosted Model	PCA + Remove Outliers	0.856	0.712	0.884	0.736	0.924	0.900	0.107	0.139

Estrategias de validación

Validación clínica

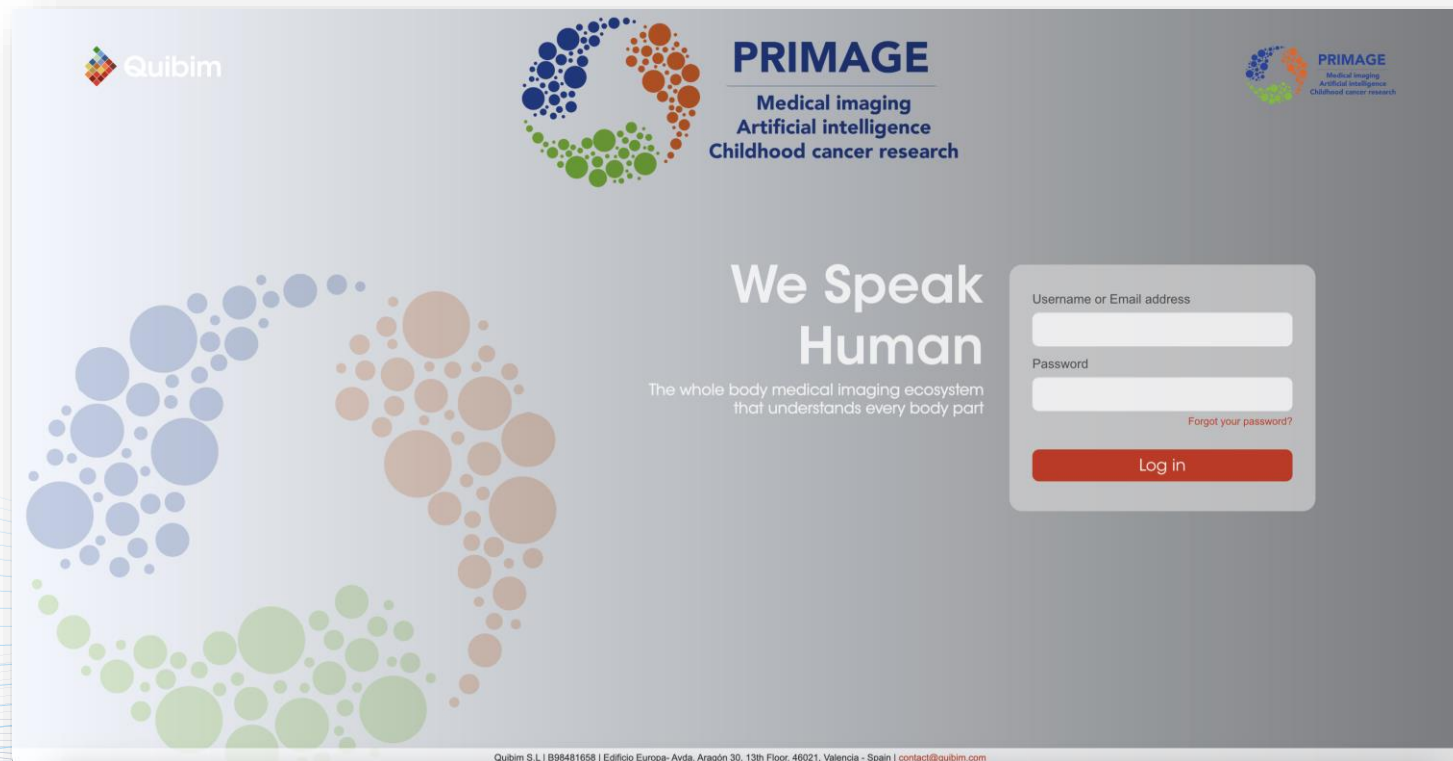
Validación de un algoritmo para la predicción del tiempo de supervivencia en pacientes con neuroblastoma



Índice

- Creación de la base de datos
- Anotación
- Estrategias de entrenamiento:
 - Modelos basados en características
 - Modelos basados en imagen
- Estrategias de validación
- **Despliegue de modelos de IA**

Despliegue de modelos de IA



The image shows a screenshot of the PRIMAGE website. The page has a light blue and grey background with a large graphic of a human figure composed of colored dots (blue, orange, green) on the left. The PRIMAGE logo is in the top right, and the Quibim logo is in the top left. The main heading is 'We Speak Human' with the tagline 'The whole body medical imaging ecosystem that understands every body part'. On the right, there is a login form with fields for 'Username or Email address' and 'Password', a 'Forgot your password?' link, and a red 'Log in' button. The footer contains contact information for Quibim S.L.

Quibim

PRIMAGE
Medical imaging
Artificial intelligence
Childhood cancer research

PRIMAGE
Medical imaging
Artificial intelligence
Childhood cancer research

We Speak Human

The whole body medical imaging ecosystem that understands every body part

Username or Email address

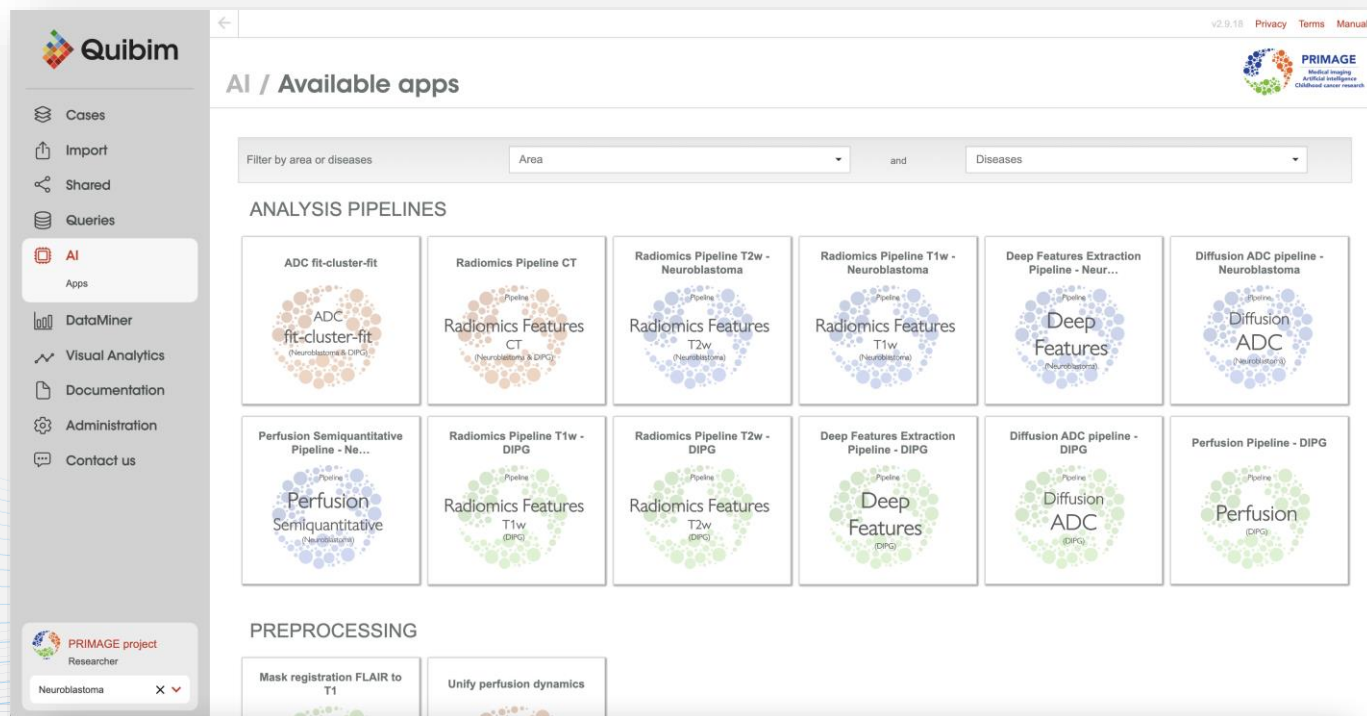
Password

[Forgot your password?](#)

Log in

























Quibim S.L. | 898481658 | Edificio Europa- Avda. Aragón 30, 13th Floor. 46021, Valencia - Spain | contact@quibim.com

Despliegue de modelos de IA



The screenshot displays the Quibim AI platform interface. On the left is a sidebar with navigation options: Cases, Import, Shared, Queries, AI (selected), DataMiner, Visual Analytics, Documentation, Administration, and Contact us. The main content area is titled "AI / Available apps" and includes a filter section with "Area" and "Diseases" dropdown menus. Below this, the "ANALYSIS PIPELINES" section features a grid of 12 pipeline cards, each with a circular icon and text: "ADC fit-cluster-fit (Neuroblastoma & DIPG)", "Radiomics Pipeline CT (Neuroblastoma & DIPG)", "Radiomics Pipeline T2w - Neuroblastoma (Neuroblastoma)", "Radiomics Pipeline T1w - Neuroblastoma (Neuroblastoma)", "Deep Features Extraction Pipeline - Neur... (Neuroblastoma)", "Diffusion ADC pipeline - Neuroblastoma (Neuroblastoma)", "Perfusion Semiquantitative Pipeline - Ne... (Neuroblastoma)", "Radiomics Pipeline T1w - DIPG (DIPG)", "Radiomics Pipeline T2w - DIPG (DIPG)", "Deep Features Extraction Pipeline - DIPG (DIPG)", "Diffusion ADC pipeline - DIPG (DIPG)", and "Perfusion Pipeline - DIPG (DIPG)". The "PREPROCESSING" section at the bottom shows two cards: "Mask registration FLAIR to T1" and "Unify perfusion dynamics". The top right of the interface shows the version "v2.9.18" and links for "Privacy", "Terms", and "Manual". A "PRIMAGE" logo is also present in the top right corner.

Despliegue de modelos de IA

VIBE-IPATTARDIO	Images: 56	  
VIBE-IPATVACIO	Images: 56	  
VIBE-IPATVACIO	Images: 56	  
VIBE-IPATVENOSO	Images: 56	  
VIBE-IPATVENOSO	Images: 56	  
T2W_Preprocessed	Images: 26	  
ADC_map_Mask1	Images: 24	  
DCE_1800	Images: 336	  

Report / Radiomics Pipeline T2w - Neuroblastoma

[Back to cases](#)

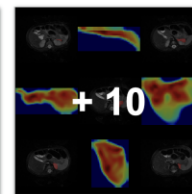
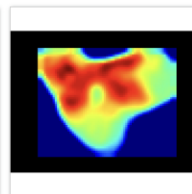
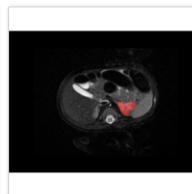
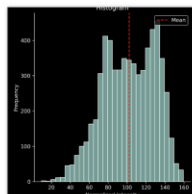
Study results

[Download all](#) [Download csv](#) [Download xls](#)

Actions	ROI ID	ROI	Patient	Patient ID	Shape - Elongation Value	Shape - Flatness Value	Shape - Least axis length Value	Shape - Major axis length Value	Shape - Maxia
	c9f2219468b9dc2dbfa4c7fd6965056		01_Neuroblastoma_6	01_Neuroblastoma_6	0.649485	0.461158	13.986489	30.329061	33.941125

Report images

ROI: Mask1



Conclusiones

- La colección y anotación de datos es un paso esencial para garantizar un buen rendimiento y generalización del modelo.
- Cuando se desarrolla un modelo de IA basado en imagen, podemos aproximarlos de dos formas: basado en características o basado en imagen.
- La validación externa del modelo es necesaria para garantizar la robustez y generalización del modelo.
- Hay diferentes maneras de validar un modelo que dependerá de su uso final.