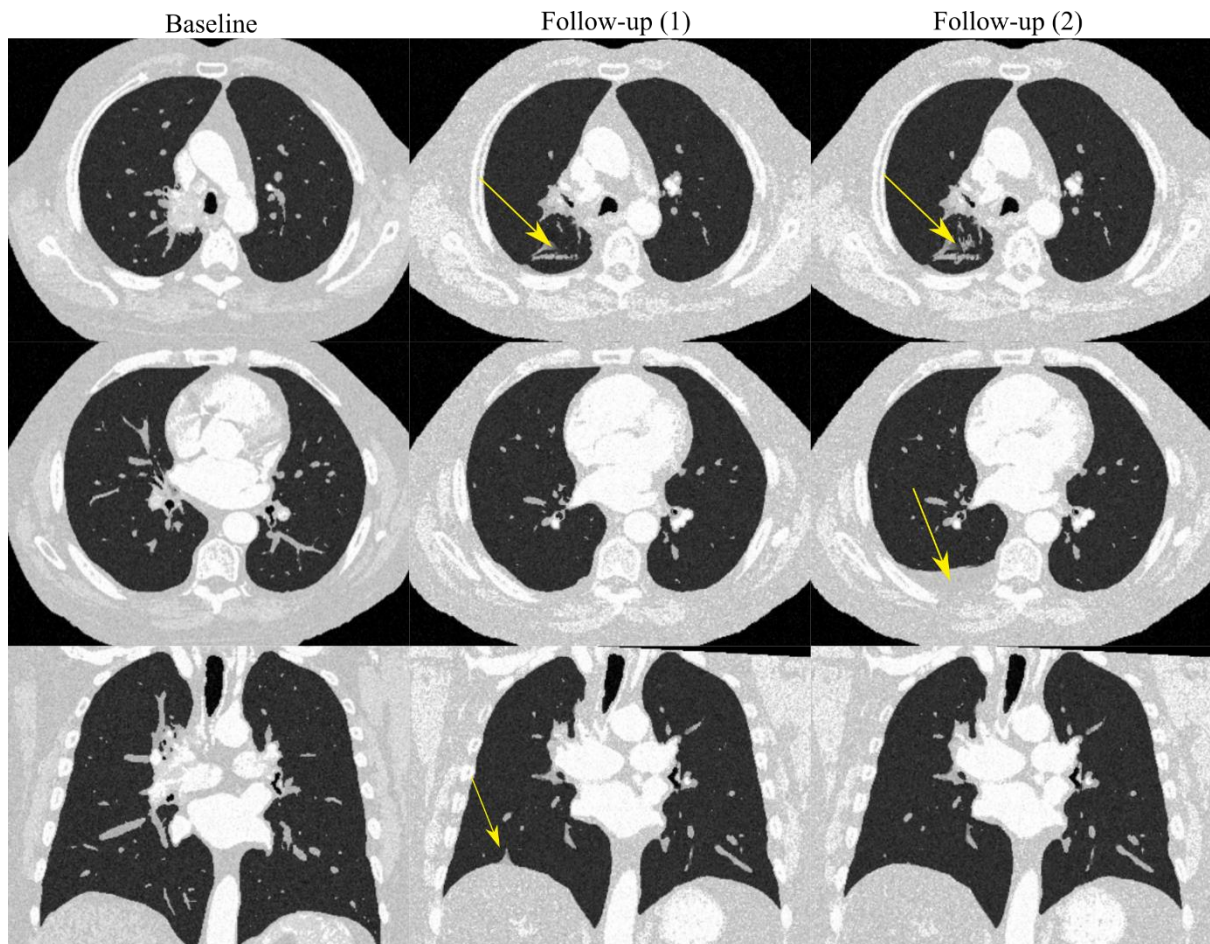


This dataset consists of two virtual phantom datasets, that represent radiation-induced lung damage (RILD) after curative RT.

The two datasets contain the same baseline CT scan, and two different images for possible follow-up. These differ in the volume of parenchymal damage, and presence/absence of pleural effusion and diaphragmatic tenting (see Figure below).



DATA DESCRIPTION

Total of two virtual patients provided: phantom001, phantom002

For each one, the following folder structure follows:

- Baseline_data: contains all information of pre-RT scan
 - segmentations:
 - Major Airways (at least up to 4th generation): airways.nii
 - Bone anatomy (to define patient specific coordinate system): bones_for_PCA.nii
 - Chest wall surface (for both ipsi and contralateral lungs): chestwallI_ipsi.nii/ chestwall_contra.nii
 - Lungs (semi-automated segmentation of left and right lung, and ipsilateral thoracic cage): lungs.nii
 - Diaphragm surface (for both ipsi and contralateral lungs): diaphragm_ipsi.nii/ chest_contra.nii
 - Normal lung volume (NV, for both ipsi and contralateral lungs): nvolume_ipsi.nii/ nvolume_contra.nii
 - Spinal canal: scanal.nii
 - transformations:
 - Affine transformation from raw to PSC coordinate system: psc_baseline.txt
 - Raw CT image: CT_baseline.nii
 - PS transformed CT image: psc_CT_baseline.nii
- Follow_up_data: contains all information of pos-RT scan
 - Raw segmentations:
 - Major Airways (at least up to 4th generation): airways.nii
 - Chest wall surface (for both ipsi and contralateral lungs): chestwallI_ipsi.nii/ chestwall_contra.nii
 - Lungs (semi-automated segmentation of left and right lung, and ipsilateral thoracic cage): lungs.nii
 - Chest wall surface (for both ipsi and contralateral lungs): chestwallII_ipsi.nii/ chestwall_contra.nii
 - Normal lung volume (NV, for both ipsi and contralateral lungs): nvolume_ipsi.nii/ nvolume_contra.nii
 - High-intensity lung volume (NV, for both ipsi and contralateral lungs): hivolume_ipsi.nii/ hivolume_contra.nii
 - Spinal canal: scanal.nii
 - Diaphragm surface (for both ipsi and contralateral lungs, defined for CT image resampled in baseline space): psc_diaphragm_ipsi_resampled.nii/ psc_contra_ipsi_resampled.nii
 - Transformations:
 - Affine transformation from raw to PSC coordinate system: psc_follow_up.txt
 - Affine transformation to baseline scan: followup2baseline.txt
 - Raw CT image: CT_followup.nii
 - PSC transformed CT image: psc_CT_followup.nii
 - CT image resampled to baseline space: psc_CT_followup_resampled.nii

BIOMARKER CALCULATION

Details of how the features are used to calculate the biomarkers can be found in the spreadsheet [here](#).

Using the segmentation data provided with the dataset, the values of the biomarkers calculated are:

phantom001:

				FEATURES MEASURED				
				Baseline		Follow-up		BIOMARKER
Feature	units	Biomarker	units	IL	CL	IL	CL	
NV	cm^3	ΔNV	%	3169	2757	2453	2368	11
RV	1	RV	1	N/A	N/A	0.027	0.026	1.0
X	mm	ΔX	%	164	156	158	160	6
Z	mm	ΔY	%	211	226	184	196	-1
h	mm	Δh	mm	11	N/A	8	N/A	-3
C	N/A	ΔC	%	0.011	0.011	0.011	0.011	5
S	mm^2	ΔS	mm^2	0	N/A	118	N/A	118
α	deg	Δα	%	127	125	125	124	0
M	mm	ΔM	%	118	128	113	125	2
β	deg	Δβ	deg	1	N/A	3	N/A	2
t	mm	Δt	1	5	N/A	8	N/A	1.6
P	%	ΔP	%	3	N/A	10	N/A	7

phantom002:

				FEATURES MEASURED				
				Baseline		Follow-up		BIOMARKER
Feature	units	Biomarker	units	IL	CL	IL	CL	
NV	cm^3	ΔNV	%	3169	2757	2401	2367.96	14
RV	1	RV	1	N/A	N/A	0.028	0.026	1.1
X	mm	ΔX	%	164	156	158	160	6
Z	mm	ΔY	%	211	226	184	197	0
h	mm	Δh	mm	11	N/A	6	N/A	-5
C	N/A	ΔC	%	0.011	0.011	0.011	0.011	7
S	mm^2	ΔS	mm^2	0	N/A	46	N/A	46
α	deg	Δα	%	127	125	125	124	0
M	mm	ΔM	%	118	128	113	125	2
β	deg	Δβ	deg	1	N/A	3	N/A	2
t	mm	Δt	1	5	N/A	8	N/A	1.6
P	%	ΔP	%	3	N/A	14	N/A	11