

Before feature extraction, several steps were involved to preprocess images: (1) use threshold to exclude voxels that didn't have an intensity within the range of (mean-standard deviation, mean+8\*standard deviation), to remove the noise; (2) normalize images using z-score normalization with z=100; (3) resample images to  $1*1*1 \text{ mm}^3$ ; (4) discretize images by a fixed bin width of 5. Below image shows the distribution of intensity range of each patient after completing image preprocessing. The intensity range had a mean of 710 and standard deviation of 203.

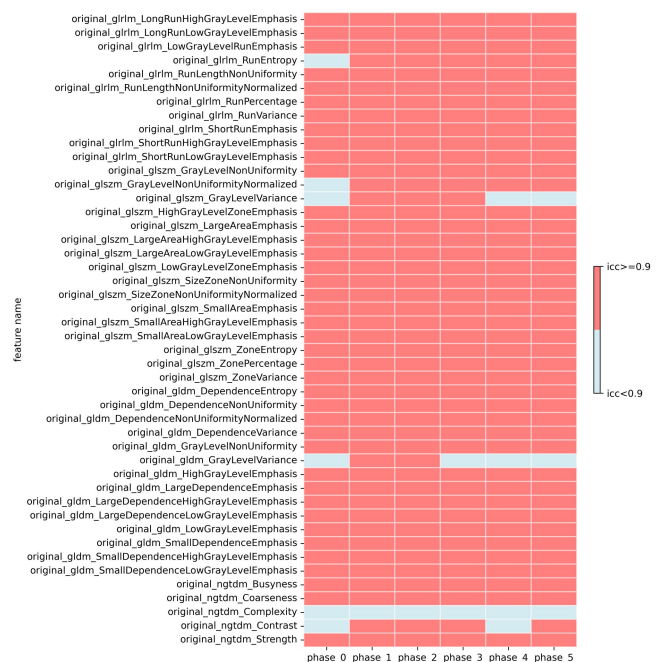


Figure S3. Feature change across different DCE-MRI phases.

The assessment of feature change across different DCE-MRI phases involved evaluating whether the feature change calculated by subtraction was significantly different from zero. It was conducted by performing a single-sample t test. A t test p value smaller than 0.05 was determined as 'change'. Below figures shown the feature change across different phases. The coordinate of x-axis n/m referred to feature change = feature at phase n – feature at phase m, and 0-5 denoted pre-contrast phase and 1<sup>st</sup> to 5<sup>th</sup> post-contrast phase.

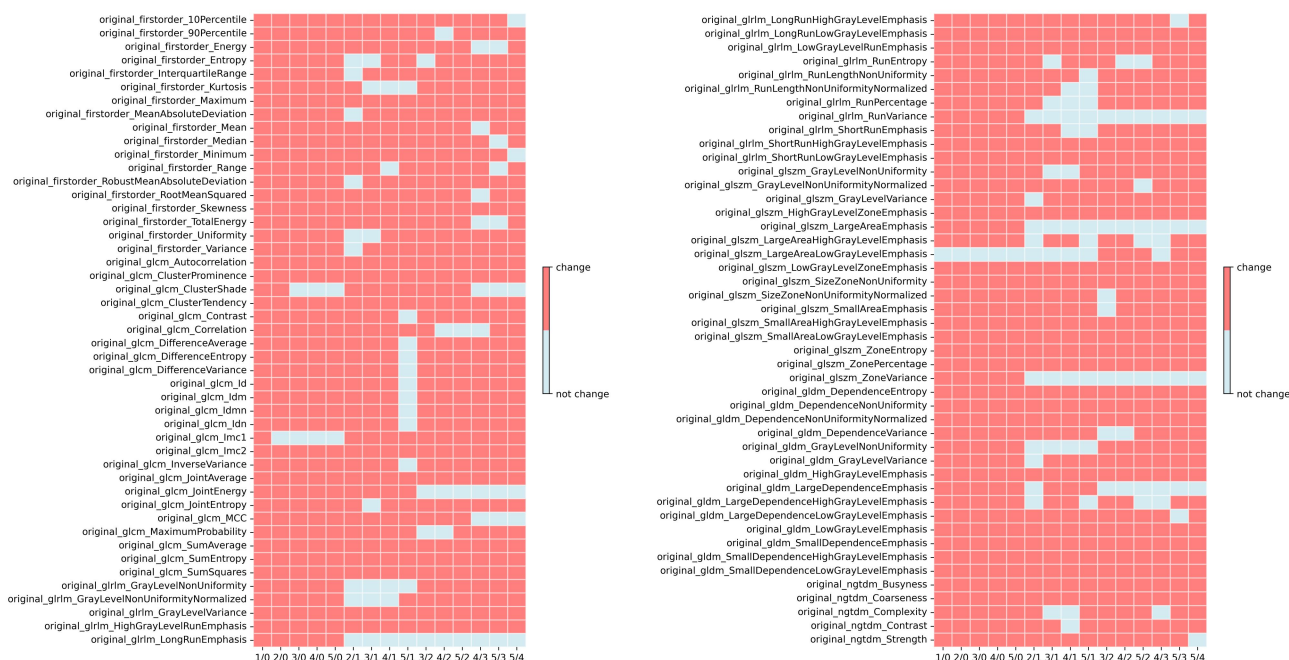


Table S1. DCE-MRI parameters

Parameter	Value
Scanner Manufacturer	GE / Philips / SIEMENS
Magnetic Field Strength	1.5T / 3.0T
Fat Suppression	Required
Slice Orientation	Axial
Slice Thickness	0.8 - 3 mm
Pixel Size	0.3 - 1.4 mm
Slice Number	56 - 256
Repetition Time	3.8 - 9.3 s
Echo Time	1.3 - 4.8 s
Number of averages	0.7 - 3
Spacing Between Slices	0.8 - 2.6 mm
Flip Angle	10 - 20 degrees
Total Phase	6 - 11
Sequence Acquisition Time	80 - 100 s
Total Acquisition Time	At least 8 min after contrast agent injection

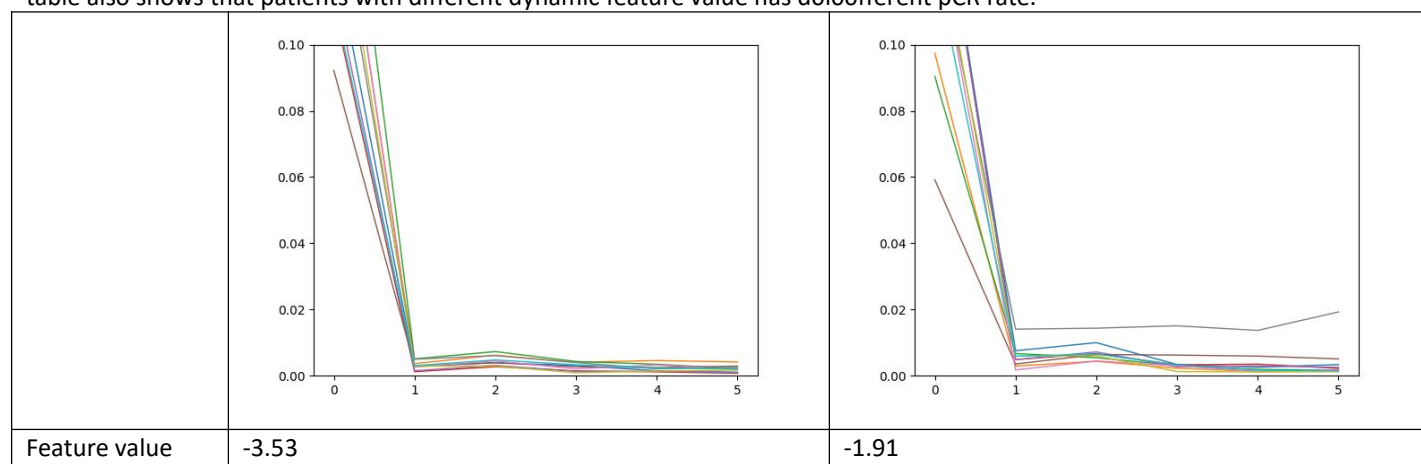
Table S2. Patient characteristics of pCR and non-pCR.

	non-pCR	pCR	P-value
<b>Treatment</b>			
Paclitaxel	114	28	<0.001
Paclitaxel + ABT 888 + Carboplatin	36	21	
Paclitaxel + AMG 386	67	29	

	Paclitaxel + AMG 386 + Trastuzumab	10	6	
	Paclitaxel + Ganetespib	58	20	
	Paclitaxel + Ganitumab	69	20	
	Paclitaxel + MK-2206	32	14	
	Paclitaxel + MK-2206 + Trastuzumab	15	15	
	Paclitaxel + Neratinib	54	31	
	Paclitaxel + Pembrolizumab	29	25	
	Paclitaxel + Pertuzumab + Trastuzumab	13	20	
	Paclitaxel + Trastuzumab	17	3	
	T-DM1 + Pertuzumab	17	22	
<b>HR</b>	negative	201	153	<0.001
	positive	330	101	
<b>HER2</b>	negative	432	169	<0.001
	positive	99	85	
<b>MP</b>	negative	312	94	<0.001
	positive	219	160	
<b>Age(mean,range)</b>		49 (23-77)	49 (25-73)	0.530
<b>Race</b>	American Indian or Alaska Native	3	1	0.791
	American Indian or Alaska Native;White	0	1	
	Asian	39	18	
	Asian;White	3	1	
	Black or African American	63	30	
	Native Hawaiian or Pacific Islander	2	2	
	Native Hawaiian or Pacific Islander;White	0	1	
	White	419	199	
	N/A	2	1	
<b>Menopausal status</b>	Premenopausal	259	111	0.291
	Perimenopausal	18	10	
	Postmenopausal	152	89	
	N/A	102	44	
<b>Ethnicity</b>	Hispanic or Latino	66	38	0.573
	Not Hispanic or Latino	464	216	
	N/A	1	0	

Table S3 – Example of visualizing dynamic features

The table gives an example of how a dynamic feature quantify the change of a radiomics feature. The selected radiomics feature is original\_glszm\_LowGrayLevelZone\_Emphasis. The dynamic feature is MD\_hrv\_classic\_pnn40. It calculates the proportion of differences magnitude that is greater than 4% of standard deviation of the time series. In these figures, the x-axis denotes the phase of DCE-MRI and y-axis denotes the value of radiomics feature. And each line denotes one patient. Those have a relatively constant radiomics feature value after contrast agent injection (phase1 to phase5) were given lower dynamic feature value. This table also shows that patients with different dynamic feature value has different pCR rate.



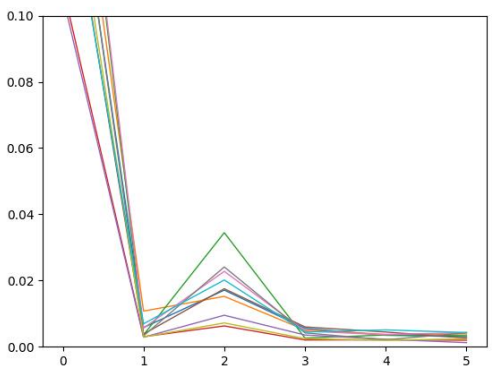
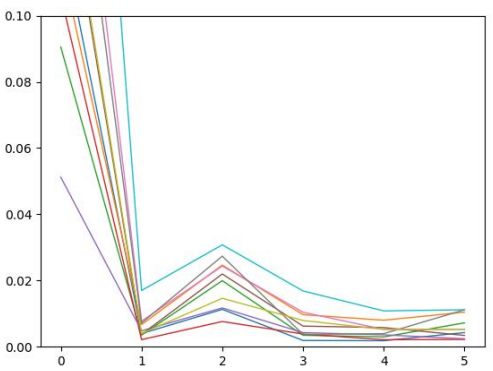
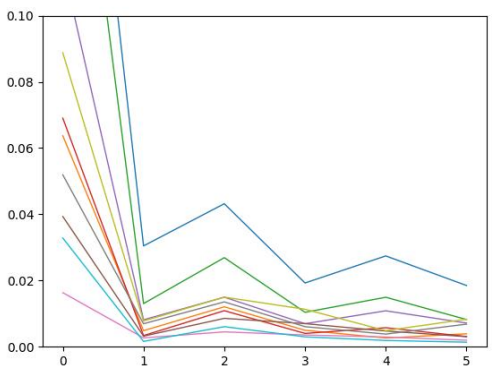
pCR rate (pCR / total)	0.2 (2/10)	0.25 (8/32)
		
Feature value	-0.29	1.33
pCR rate (pCR / total)	0.32 (185/582)	0.38 (50/133)
		
Feature value	2.95	
pCR rate (pCR / total)	0.30 (9/30)	

Table S4 Performance of models with different feature number.

<u>Dynamic</u>	<u>model</u>	<u>Training AUC</u>	<u>Internal Validation</u>	<u>Radiomic</u>	<u>model</u>	<u>Training AUC</u>	<u>Internal Validation</u>
<u>feature No.</u>			<u>AUC</u>	<u>feature No.</u>			<u>AUC</u>
<u>1</u>		<u>0.571</u>	<u>0.560</u>	<u>1</u>		<u>0.595</u>	<u>0.564</u>
<u>2</u>		<u>0.600</u>	<u>0.634</u>	<u>2</u>		<u>0.621</u>	<u>0.535</u>
<u>3</u>		<u>0.607</u>	<u>0.635</u>	<u>3</u>		<u>0.650</u>	<u>0.637</u>
<u>4</u>		<u>0.643</u>	<u>0.649</u>	<u>4</u>		<b><u>0.650</u></b>	<b><u>0.640</u></b>
<u>5</u>		<u>0.650</u>	<u>0.675</u>	<u>5</u>		<u>0.650</u>	<u>0.634</u>
<u>6</u>		<u>0.660</u>	<u>0.643</u>	<u>6</u>		<u>0.650</u>	<u>0.635</u>
<u>7</u>		<u>0.664</u>	<u>0.673</u>	<u>7</u>		<u>0.666</u>	<u>0.603</u>
<u>8</u>		<u>0.675</u>	<u>0.667</u>	<u>8</u>		<u>0.672</u>	<u>0.603</u>
<u>9</u>		<u>0.682</u>	<u>0.660</u>	<u>9</u>		<u>0.673</u>	<u>0.589</u>
<b><u>10</u></b>		<b><u>0.688</u></b>	<b><u>0.680</u></b>	<u>10</u>		<u>0.690</u>	<u>0.570</u>
<u>11*</u>		<u>0.698</u>	<u>0.651</u>				

\*As the 10-feature dynamic model achieved the highest internal validation AUC. We further explored whether adding more dynamic features can obtain better performance. Therefore, the results of 11-feature dynamic model is also listed here.

Table S5 Odds ratios showing the significance of dynamic features in the dynamic model.

<u>Dynamic feature name</u>	<u>Odds ratio</u>
<u>original firstorder RootMeanSquared SP Summaries welch rect centroid</u>	<u>1.098</u>
<u>original firstorder Mean FC LocalSimple mean3 stderr</u>	<u>0.795</u>
<u>original glrlm ShortRunLowGrayLevelEmphasis SP Summaries welch rect centroid</u>	<u>0.807</u>
<u>original glszm GrayLevelNonUniformity CO HistogramAMI even 2 5</u>	<u>0.780</u>
<u>original glszm LowGrayLevelZoneEmphasis MD hrv classic_pnn40</u>	<u>1.161</u>
<u>original glcm Id SB MotifThree quantile hh</u>	<u>0.793</u>
<u>original glcm DifferenceEntropy CO HistogramAMI even 2 5</u>	<u>1.193</u>
<u>original firstorder 90Percentile FC LocalSimple mean1 tauresrat</u>	<u>1.248</u>
<u>original glcm lmc1 CO f1ecac</u>	<u>0.868</u>
<u>original firstorder InterquartileRange DN HistogramMode 10</u>	<u>0.833</u>

Table S6 Odds ratios showing the significance of radiomic features in the radiomic model.

<u>Radiomic feature name</u>	<u>Odds ratio</u>
<u>original_shape Maximum2DDiameterColumn_phase0</u>	<u>0.793</u>
<u>original glcm lmc2_phase1</u>	<u>1.345</u>
<u>original glszm ZoneEntropy_phase0</u>	<u>0.772</u>
<u>original ngtdm Busyness_phase1</u>	<u>0.973</u>

Table S7 Coefficients and p value of variables in the CRD model

<u>Variable</u>	<u>coefficient</u>	<u>P value</u>
<u>HR</u>	<u>-1.076</u>	<u>&lt;0.001</u>
<u>HER2</u>	<u>0.837</u>	<u>&lt;0.001</u>
<u>Dynamic</u>	<u>4.879</u>	<u>&lt;0.001</u>
<u>Radiomic</u>	<u>4.169</u>	<u>&lt;0.001</u>