

# Radiogenomics Analysis Report

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-----Model inputs-----

Mode:Train  
Model:multiTaskLinearModel

Param:default  
No. of imaging features provided: 36

No. of gene features provided: 976  
SampleID check results: 'The SampleIDs match for imaging and gene features'  
No. of samples: 89

performing StandScaler normalization for imaging features  
performing StandScaler for gene features













