

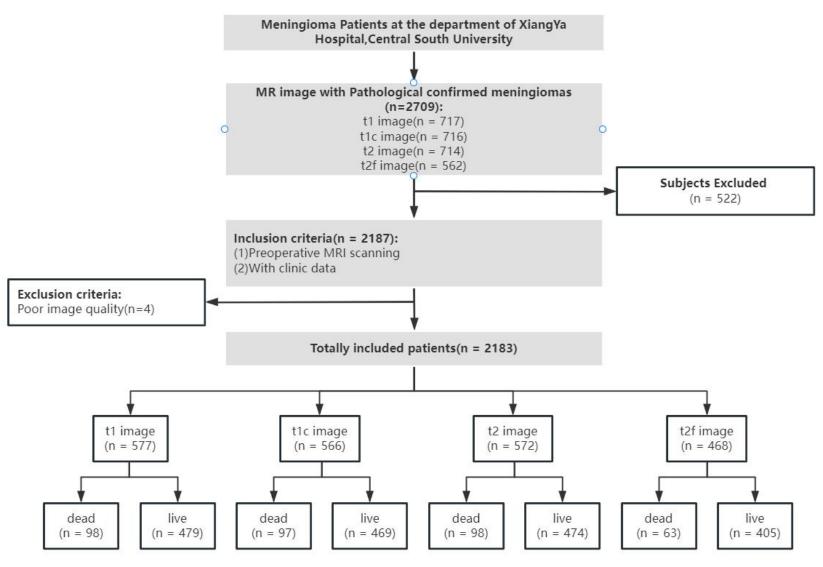
A machine learning model for predicting prognostic survival status of patients with meningiomas based on MRI radiomics and clinical features

付钰 中南大学湘雅医院药学部 2024-10-27



Sample screening process





Study framework

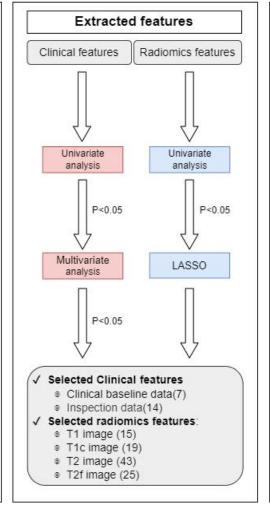
Clinical information

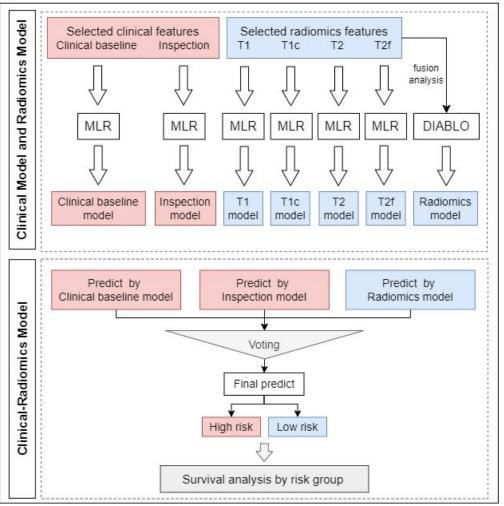


Feature Extraction Feature Selection Model Building

Clinical baseline data Preoperative PS, Extent of resection, Postoperative radiotherapy, Tumor location, Number of tumors, Tumor size, Ki-67, Age, Gender, Medication history Inspection data WBC,RBC,HGB,PLT,HCT,NE#,LY#,EO#,BA#, MO#,NE,LY,BA,EO,MO,MCV,MCH,MCHC,RDW, PCT,MPV,SG,PH,TP,ALB,GLB,A/G,TBIL,DBIL, TBA, ALT, AST, BUN, CREA, UA, K, NA, CL, Color. Clarity, BLD, LEU, GLU, KET, PRO, UBIL, UBG, NIT, VITC 60 Clinicopathological features Preoperative MRI radiomics Pyradiomics

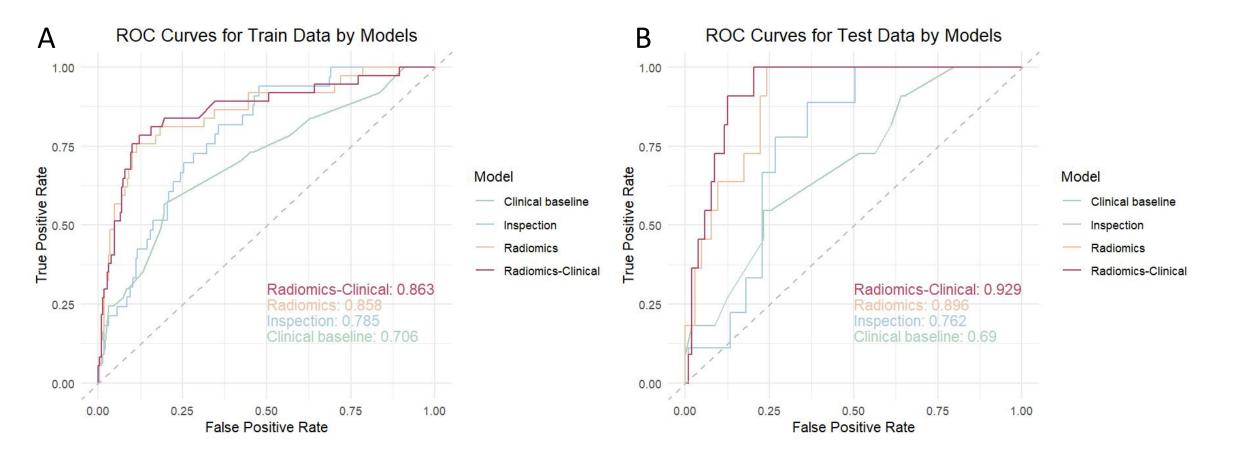
1052 radiomics features





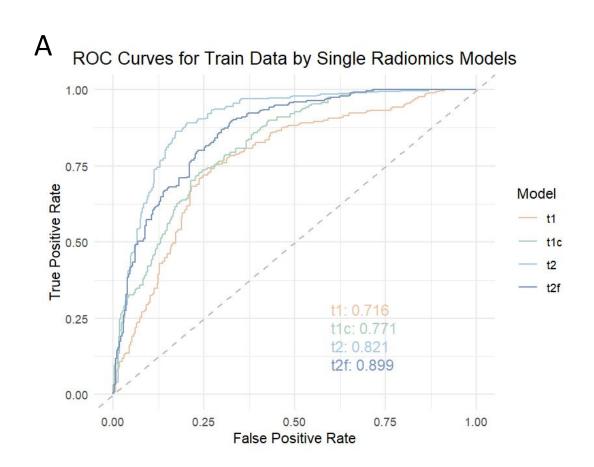
ROC Curve for Models

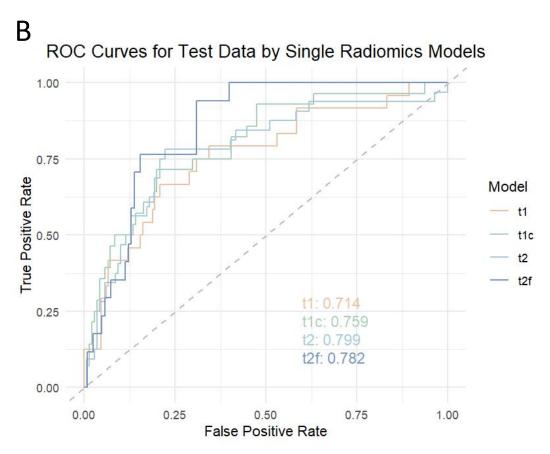




ROC Curve for Single Radiomics Models

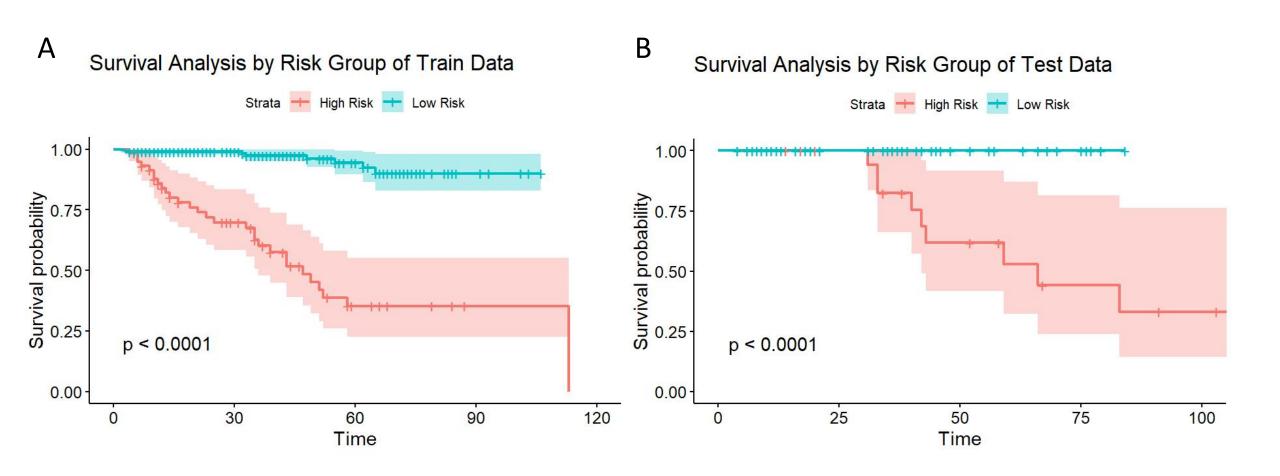






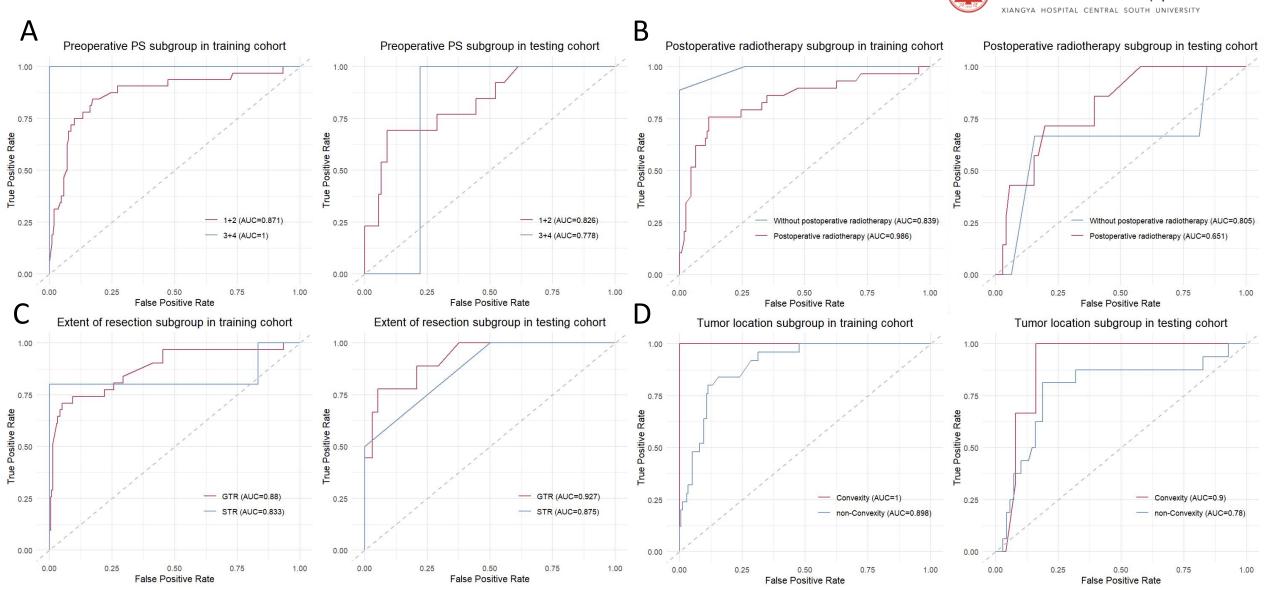
Survival Analysis by Risk Group





Subgroup analysis





Subgroup analysis



