Fraction of observed values of simulations under the model with inferred parameters Bone-Benign 1.00 -Breast-DCIS CNS-Oligo CNS-Medullo 0.75 -Kidney-ChRCC Prost-AdenoCA Bone-Epith Breast-AdenoCA Bone–Benign Kidney-RCC.clearcell sim_confint_DM_prop 0.50 -Bone-Osteosarc type Head-SCC nucleotidesubstitution1 signatures Myeloid-AML NA 0.25 -Cervix-SCC Lymph-BNHL ng-AdenoCA Melanoma.cutaneous vix-AdenoCA Kidne Stomach-Aden Co Skin-Melanoma.cutal Thy-AdenoCA 0.00 -Pancy Uterus-AdenoCA denoCA denoCA de Lung-SCC Skin-Melanoma SoftTissue-I SoftTissu Uterus-AdenoCA 0.25 0.00 0.50 0.75 1.00 -0.25sim_confint_M_prop