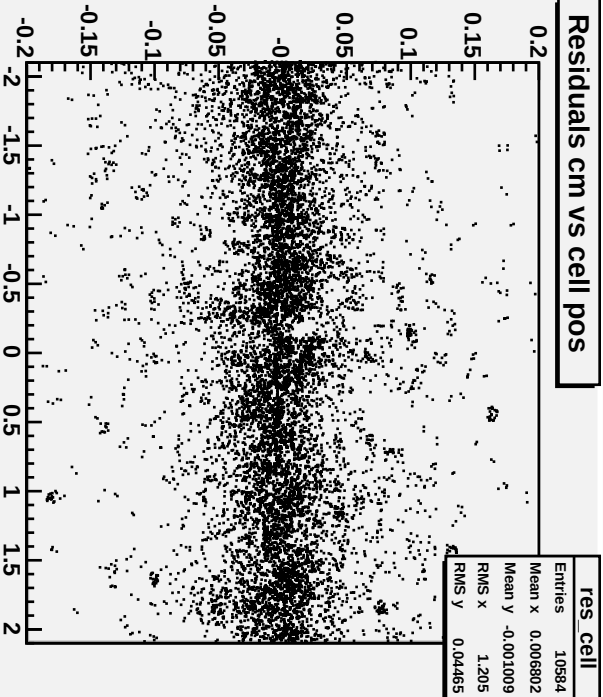
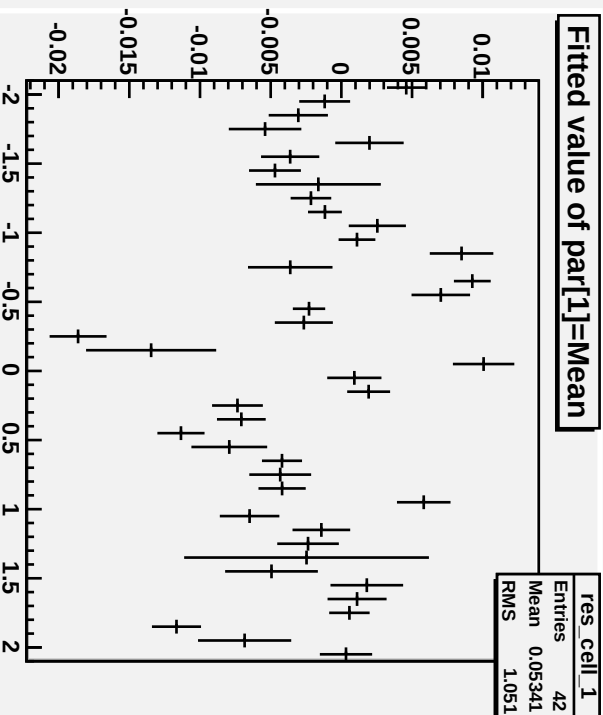


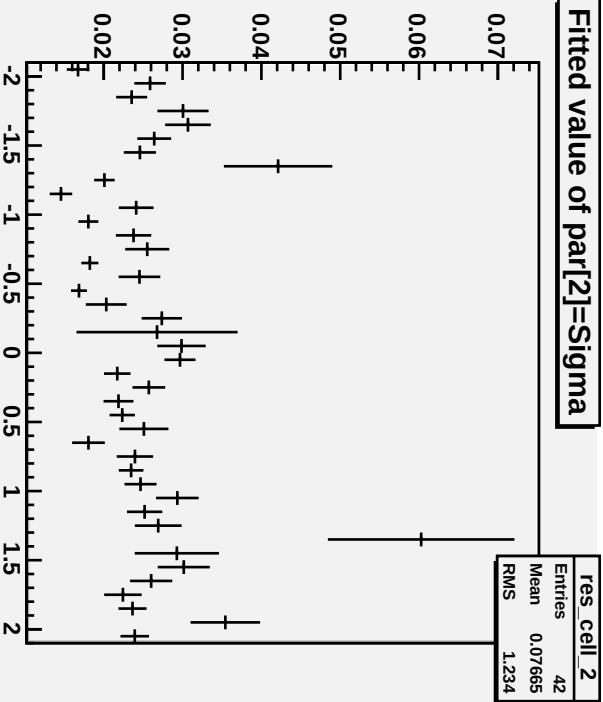
Residuals cm vs cell pos



Fitted value of par[1]=Mean



Fitted value of par[2]=Sigma



Residuals cm vs cell pos

