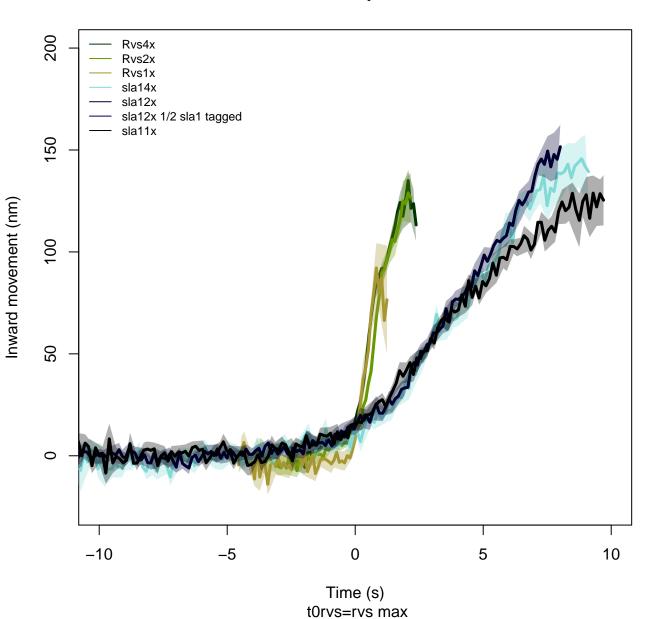
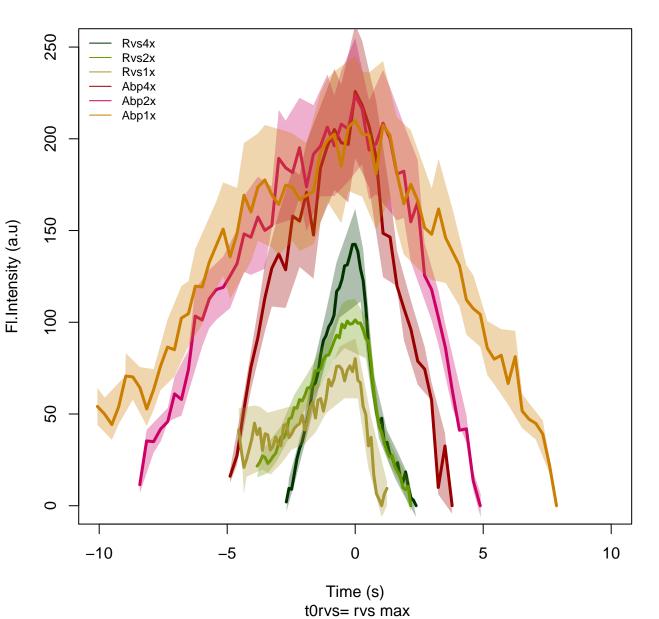
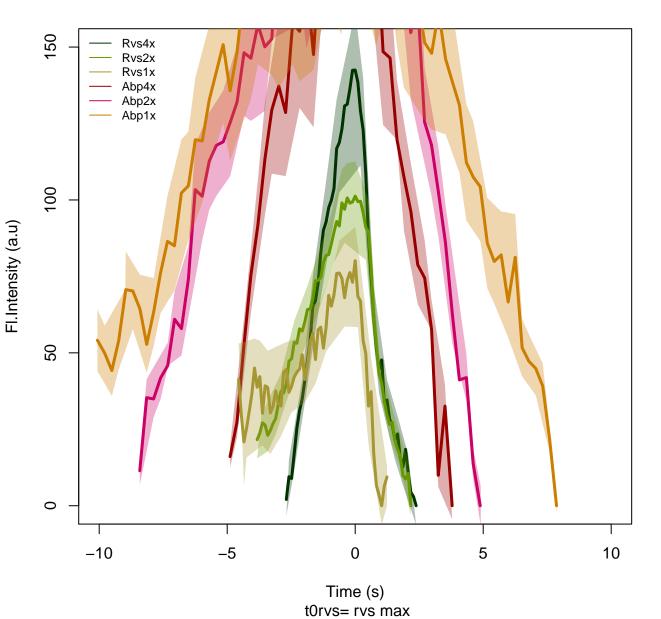
# **Rvs in diploids**



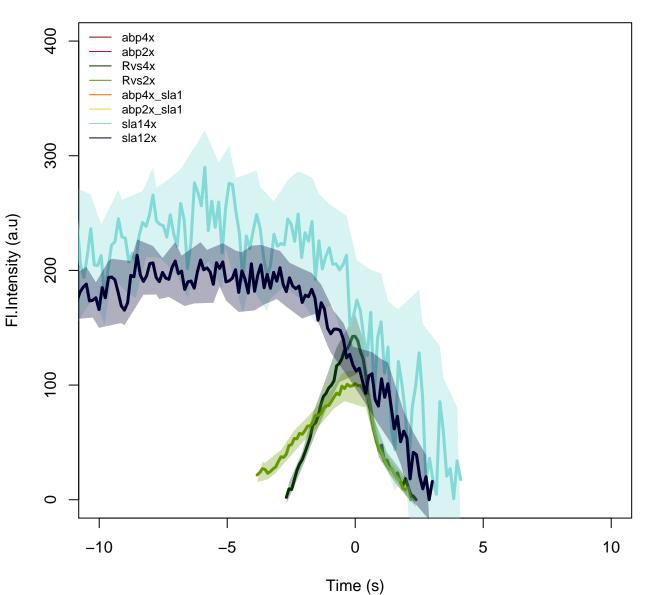
# Rvs vs Abp-Rvs



# Rvs vs Abp-Rvs

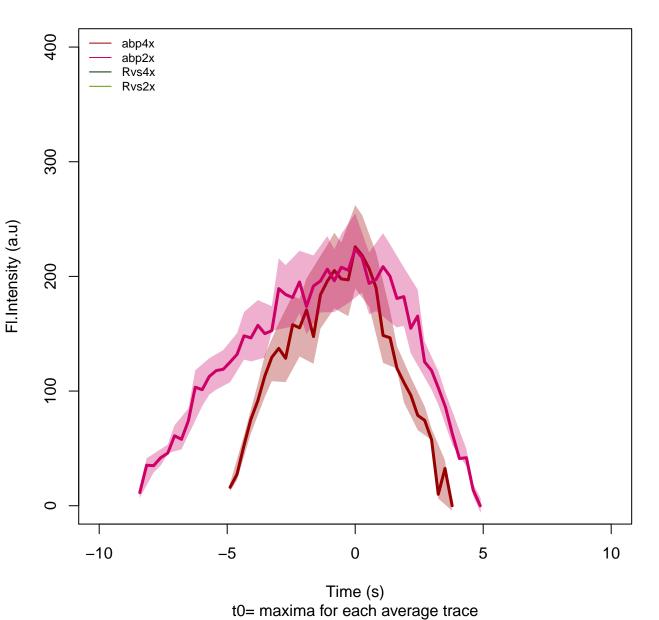


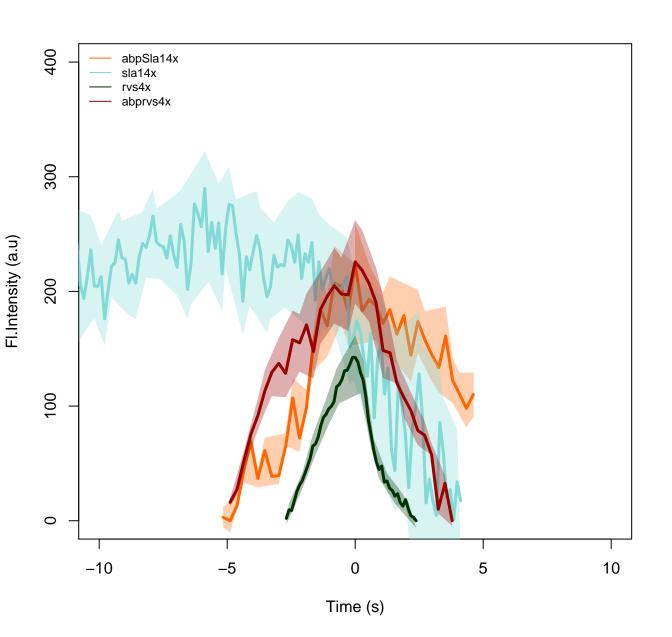
# abp1 fl.int



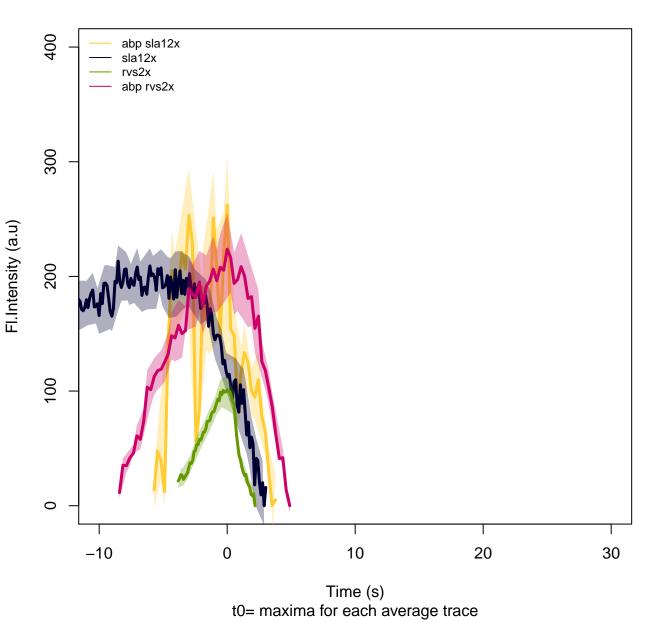
t0= maxima for each average trace

abp1 fl.int





abp1 fl.int



# abp1 in Rvs dupl diploids

