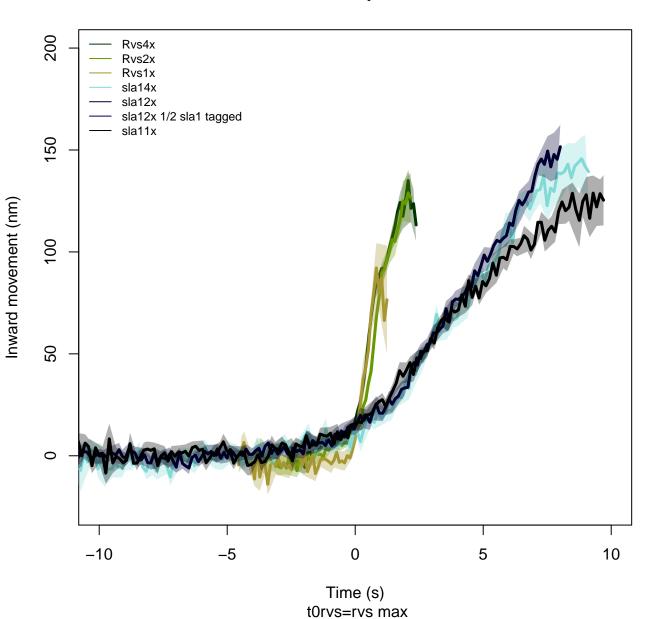
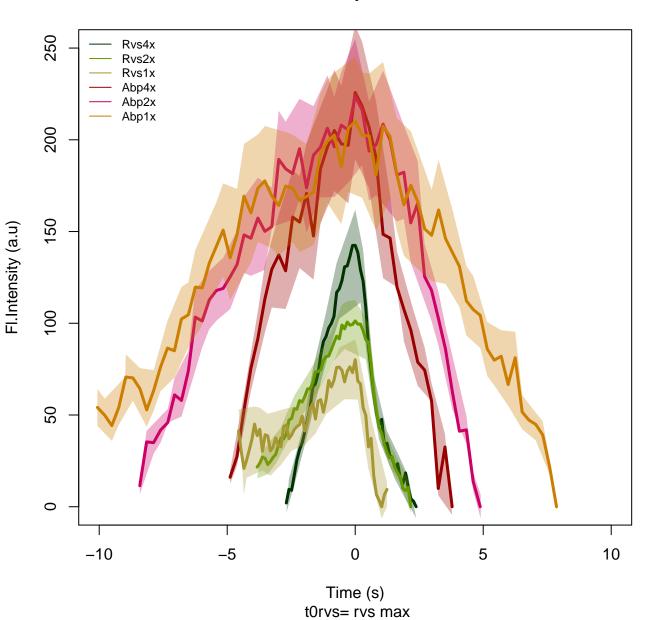
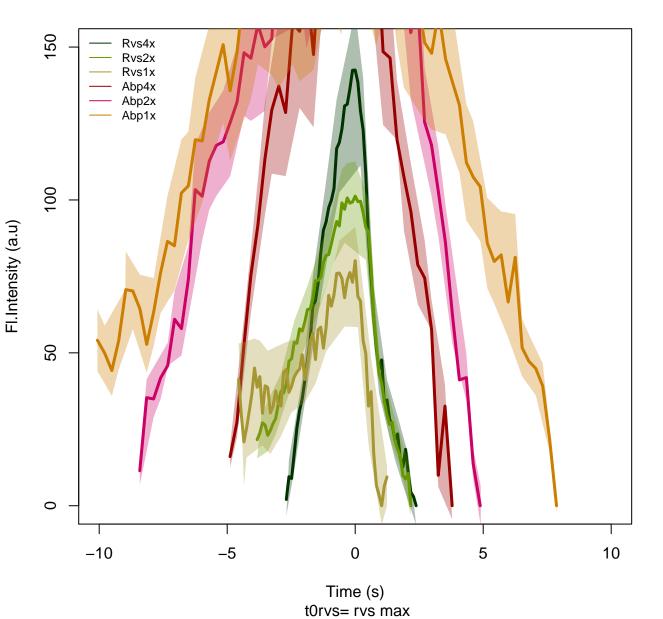
# **Rvs in diploids**



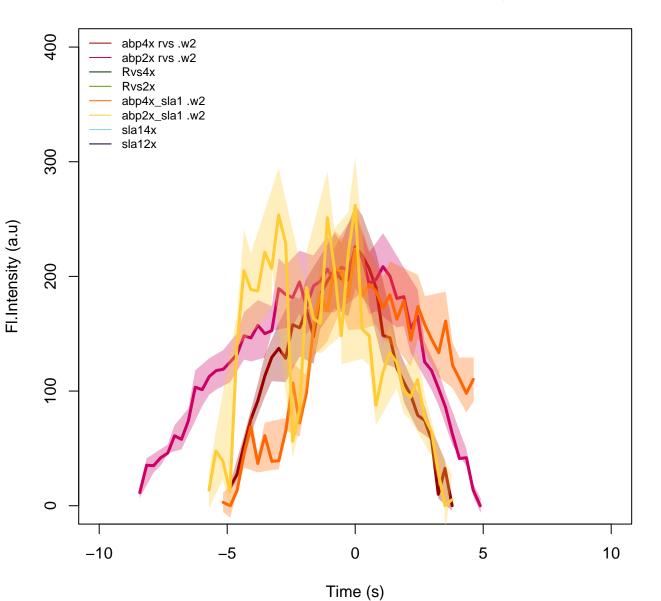
# Rvs vs Abp .w2-Rvs



# Rvs vs Abp .w2-Rvs

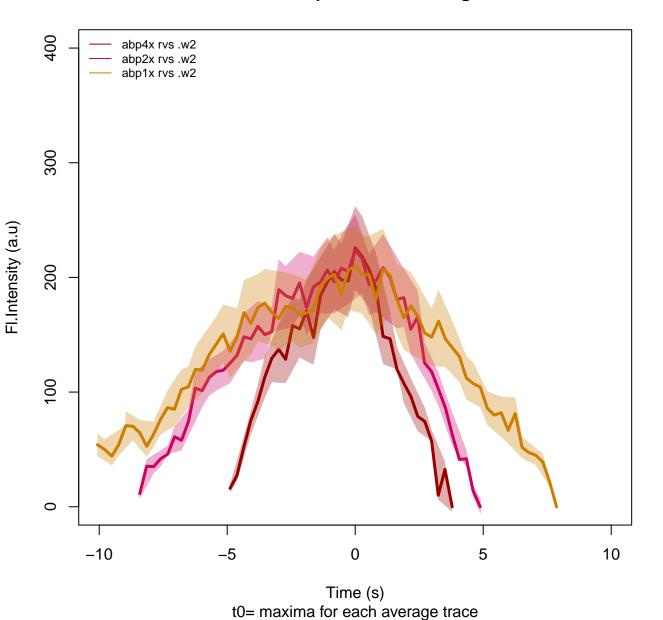


### abp1 fl.int, .w2 rvs and sla1 co-tag

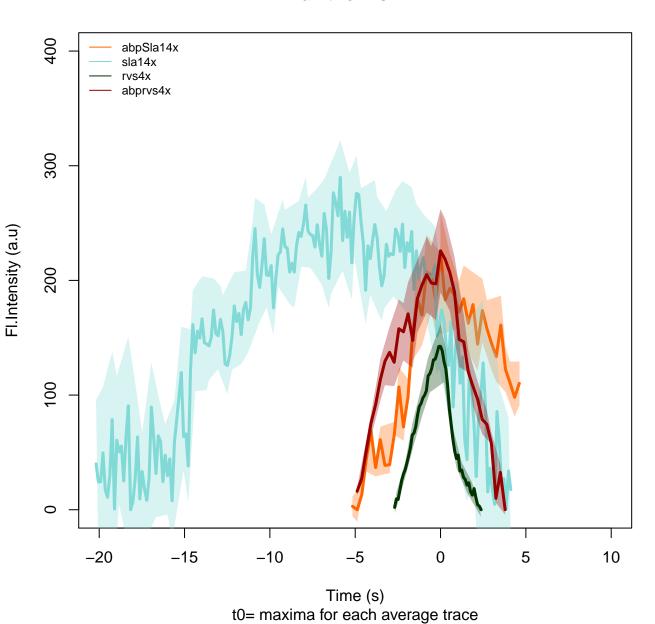


t0= maxima for each average trace

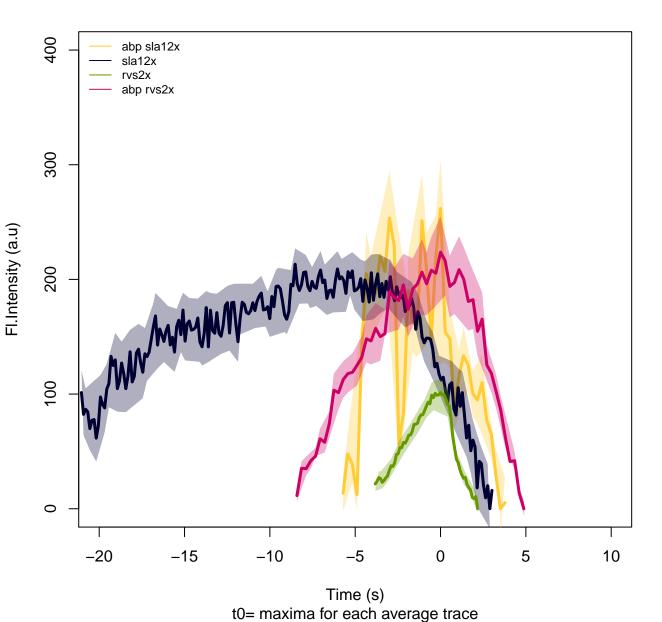
### fl. int abp1 .w2 rvs co-tag



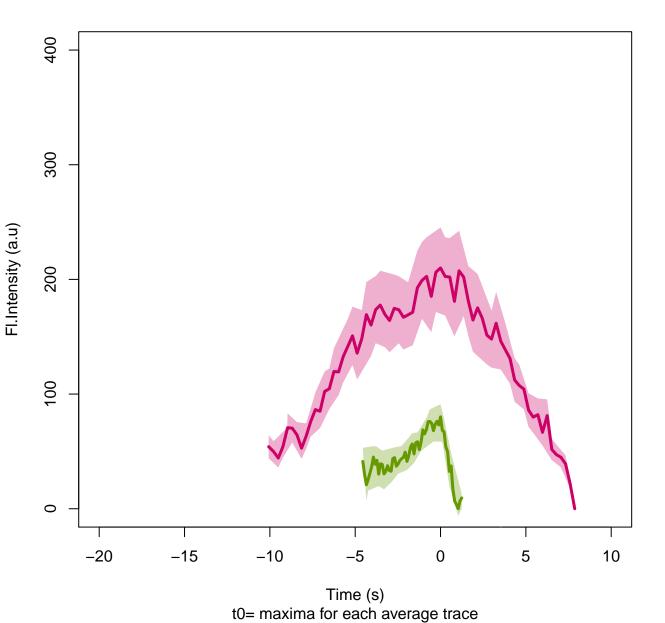
### all the rvs4x



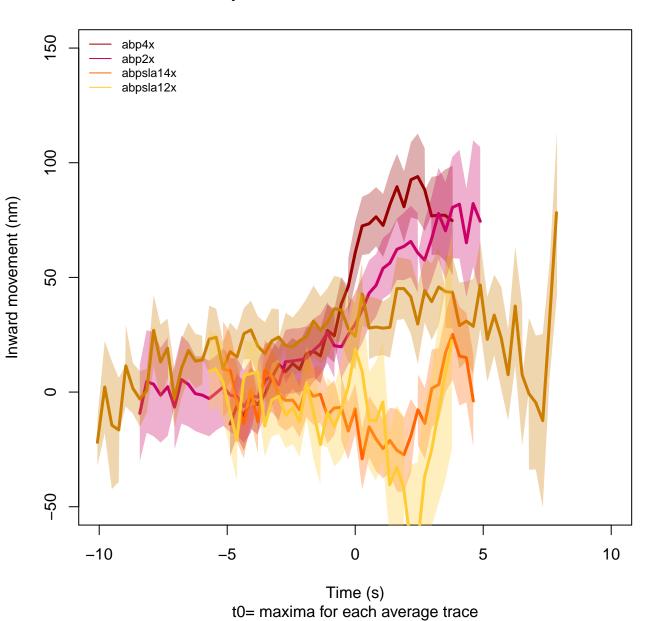
#### all the rvs2x



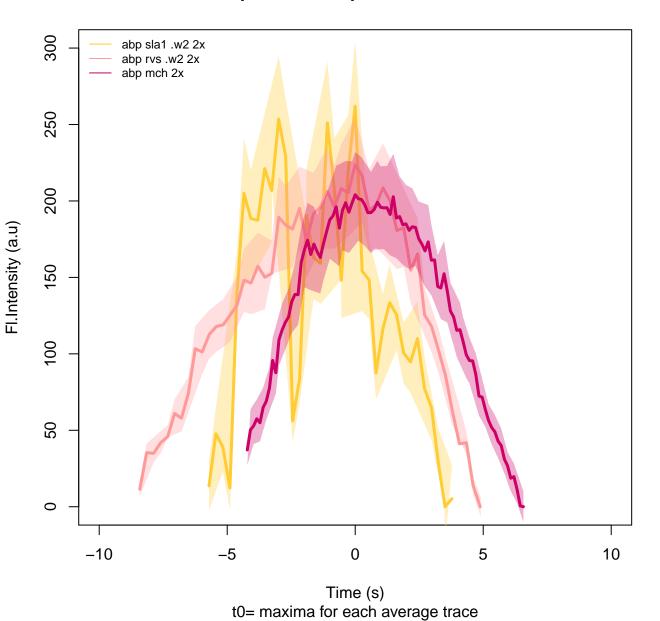
all the rvs1x

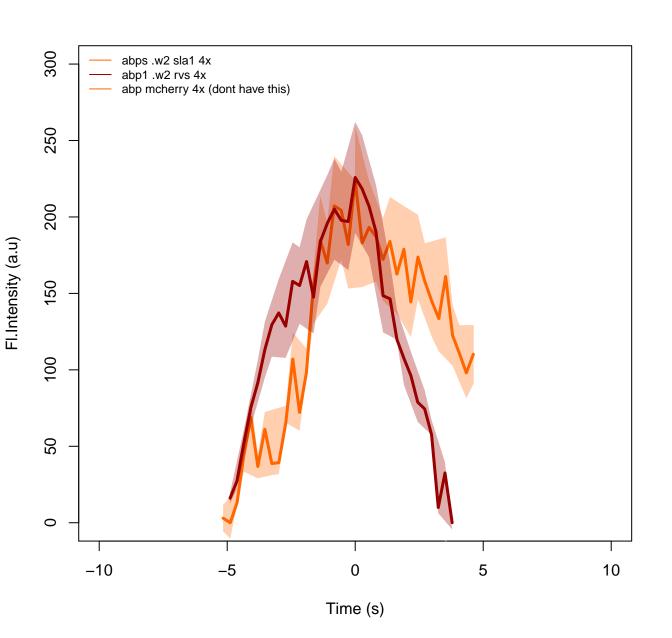


### abp1 all .w2 movements in 4x, 2x

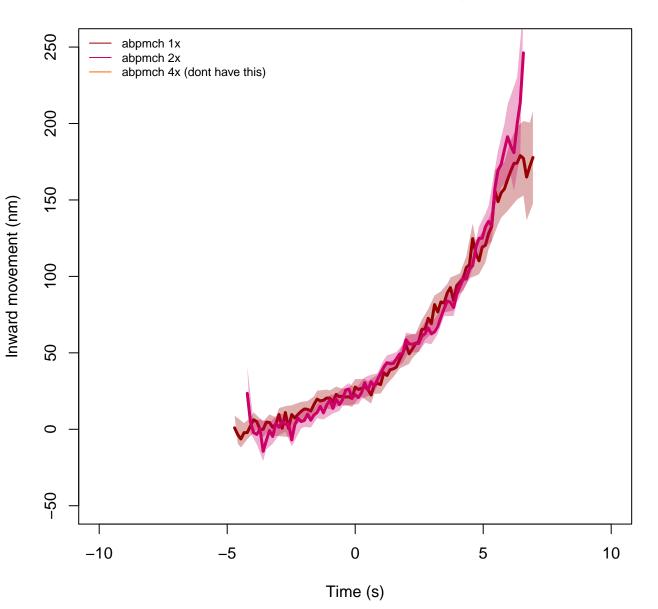


# abp1 .w2 vs abp mch in Rvs 2x





# abp1mcherry sla1 co-tag



# abp1mcherry sla1 co-tag

