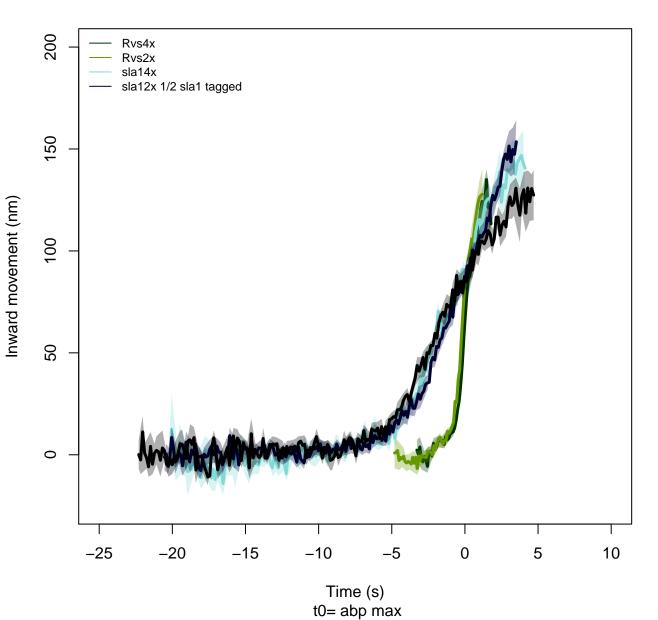
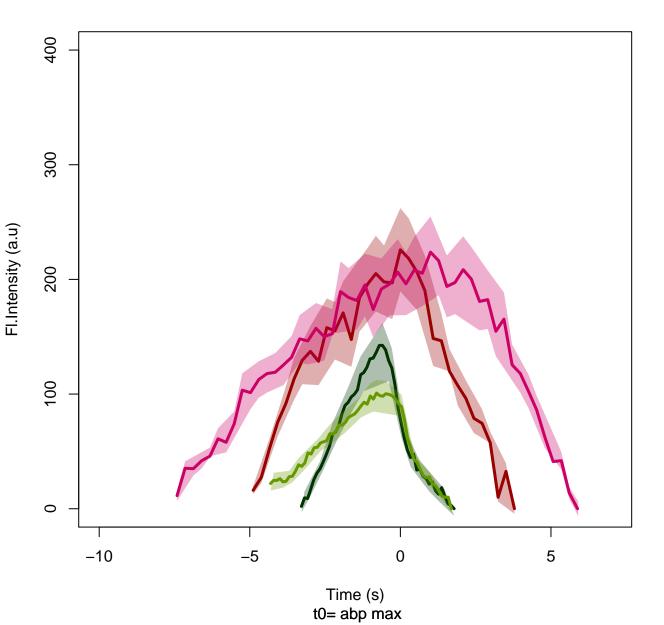
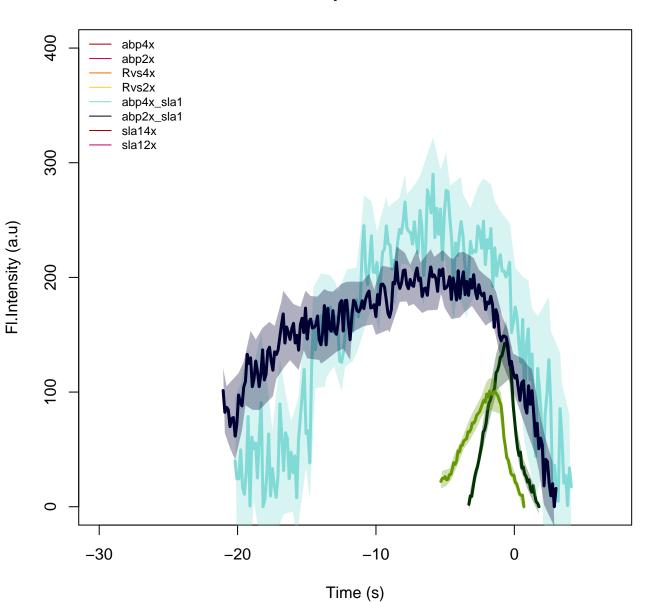
## **Rvs in diploids**



2xRvs

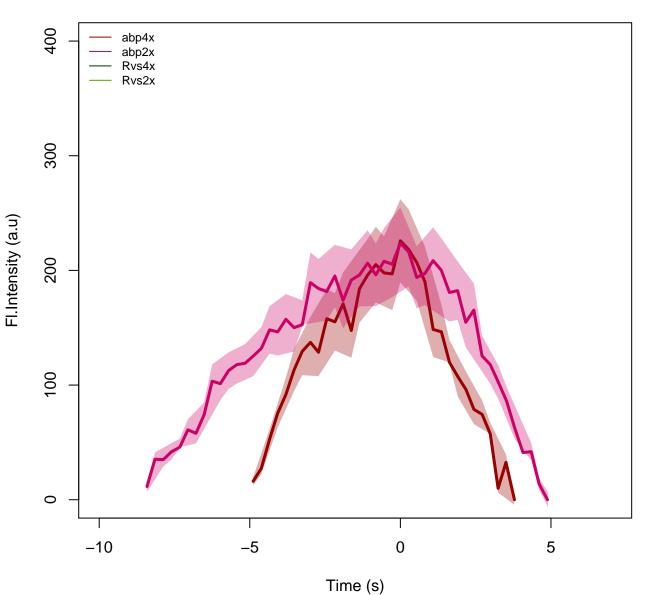


## abp1 fl.int

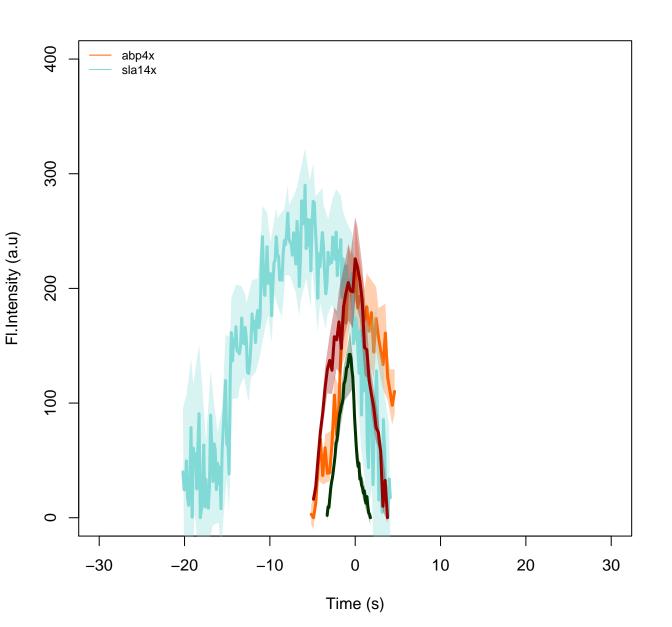


t0= maxima for each average trace

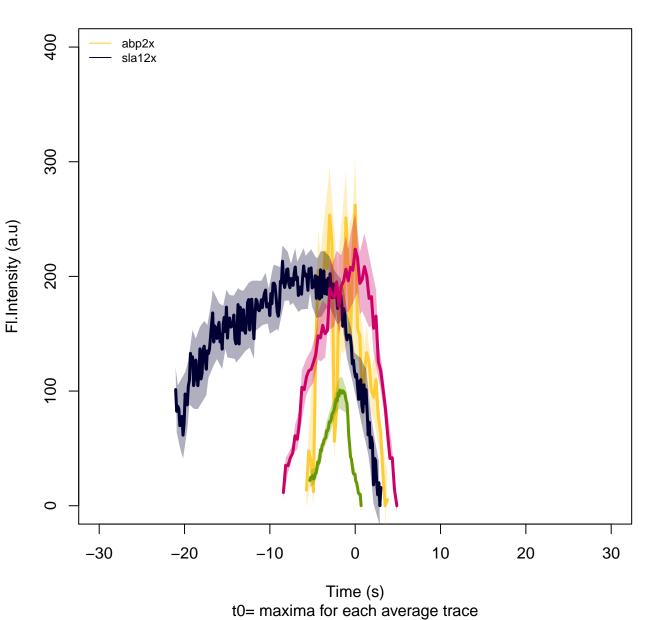
abp1 fl.int



t0= maxima for each average trace



abp1 fl.int



## abp1 in Rvs dupl diploids

