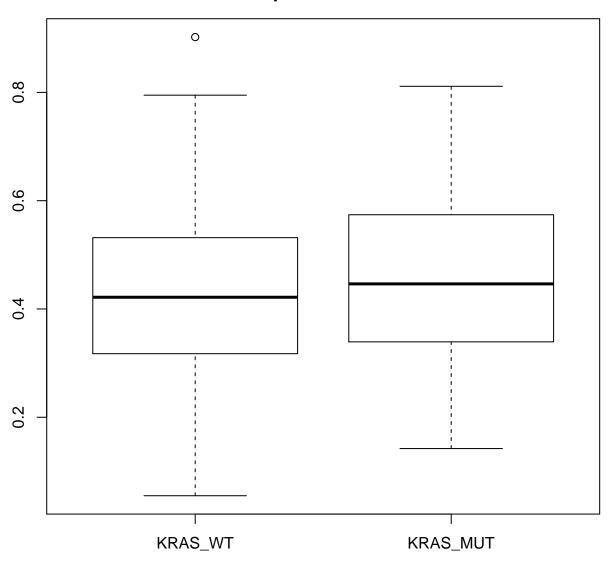
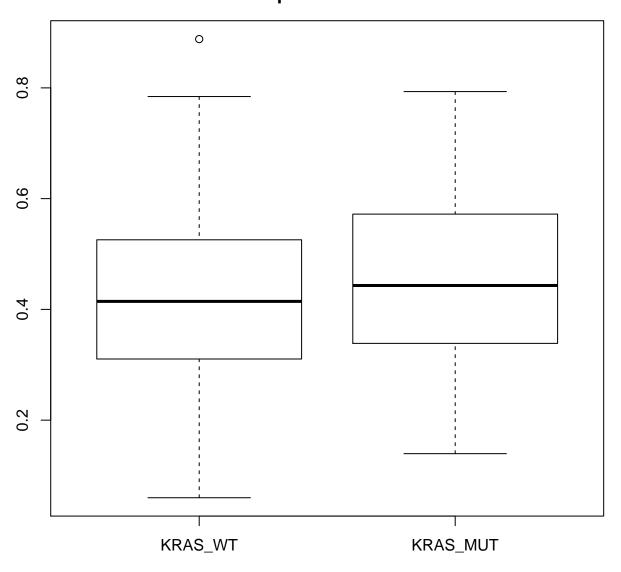
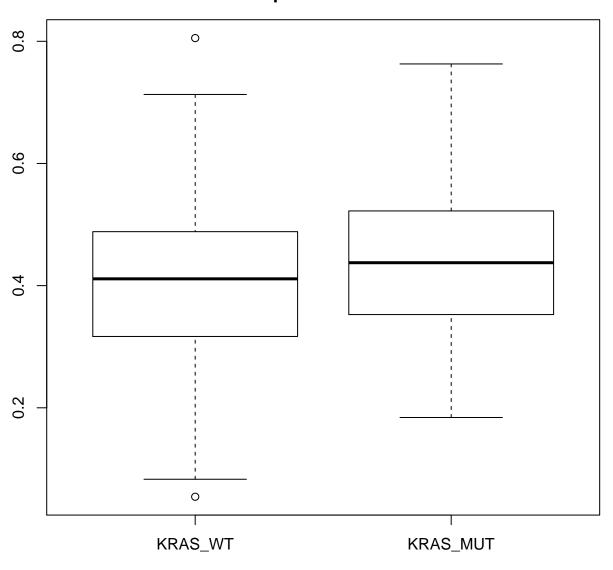
## krasgv\_100\_gene\_list/adapB\_single p-val= 0.11



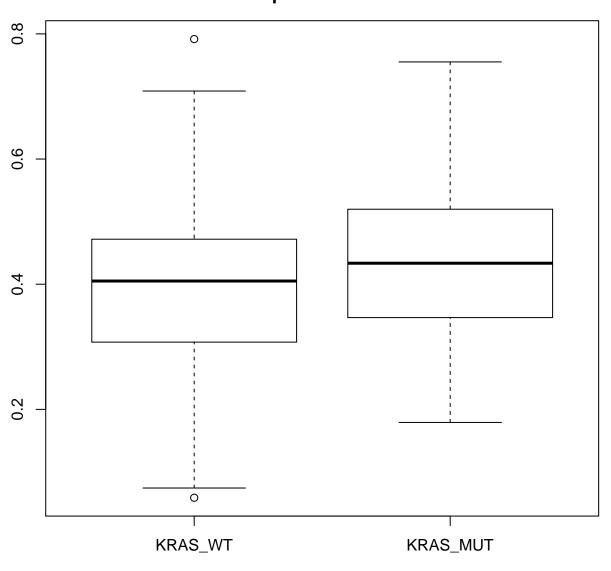
# krasgv\_100\_gene\_list/adap\_adap\_single p-val= 0.09



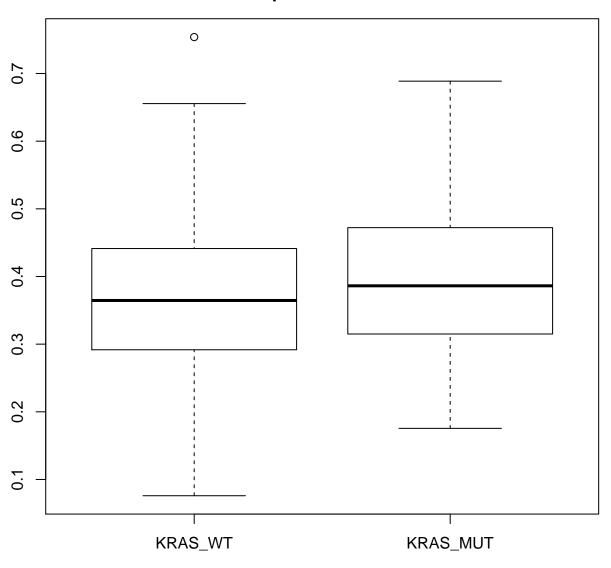
#### krasgv\_200\_gene\_list/adapB\_single p-val= 0.04



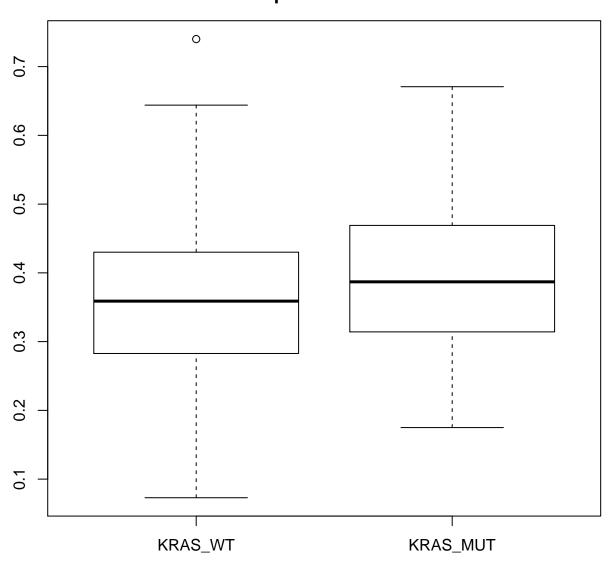
# krasgv\_200\_gene\_list/adap\_adap\_single p-val= 0.04



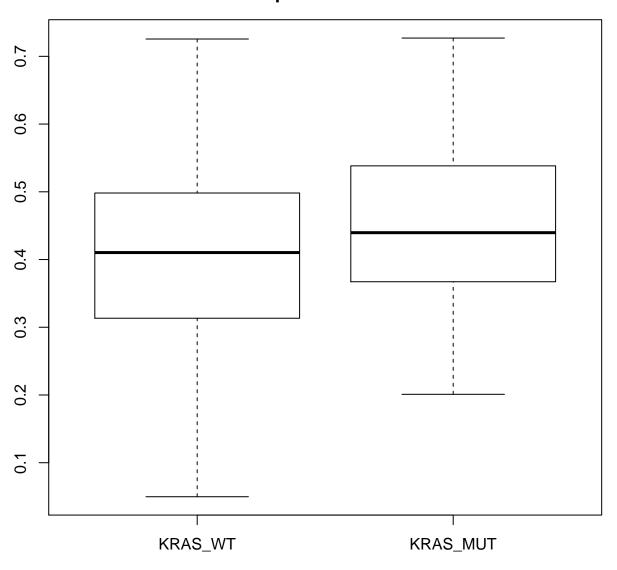
#### krasgv\_300\_gene\_list/adapB\_single p-val= 0.03



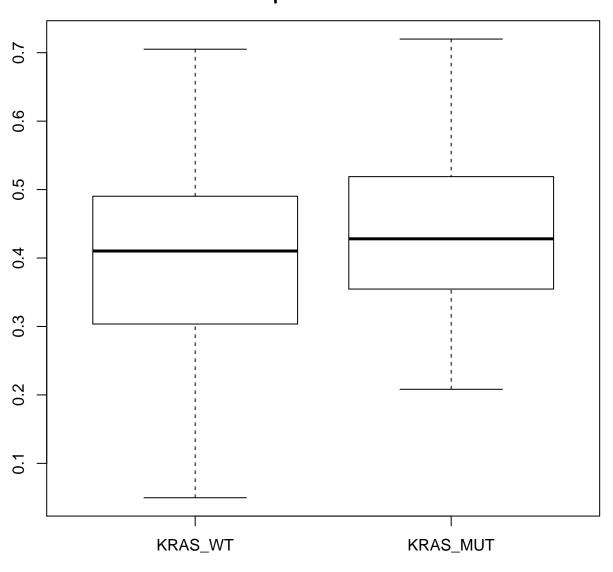
# krasgv\_300\_gene\_list/adap\_adap\_single p-val= 0.02



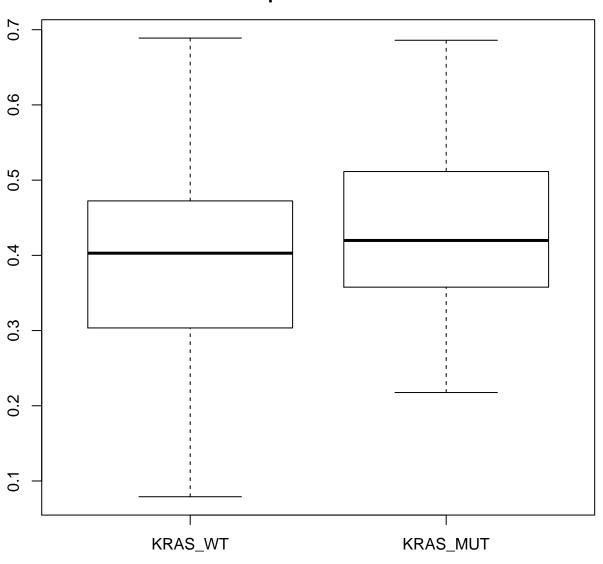
## krasqh\_100\_gene\_list/adapB\_single p-val= 0.04



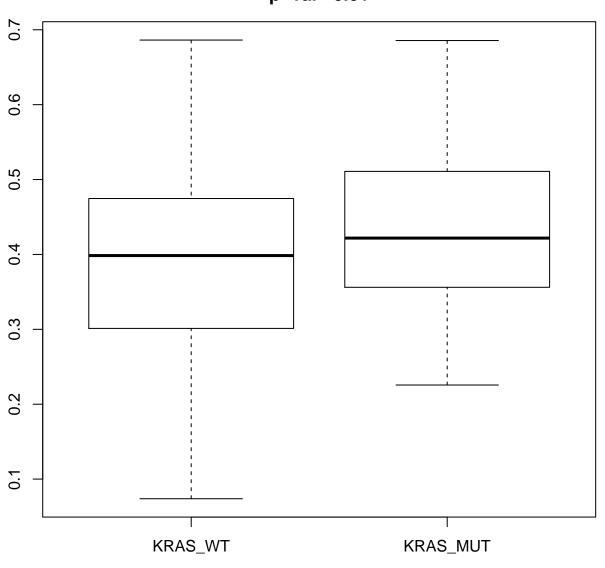
## krasqh\_100\_gene\_list/adap\_adap\_single p-val= 0.04



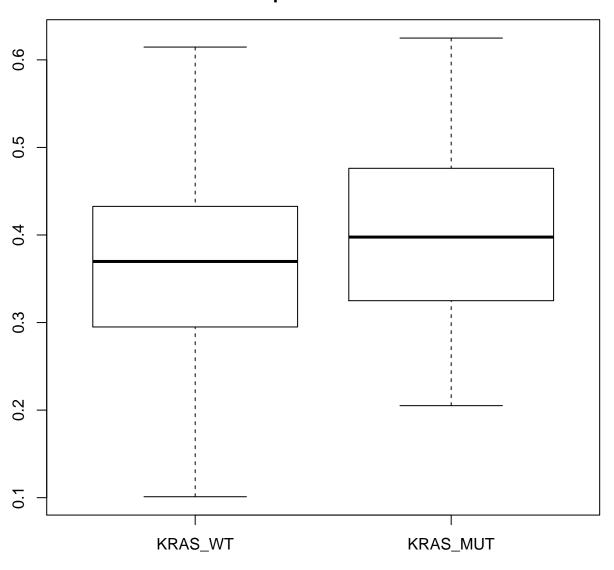
## krasqh\_200\_gene\_list/adapB\_single p-val= 0.02



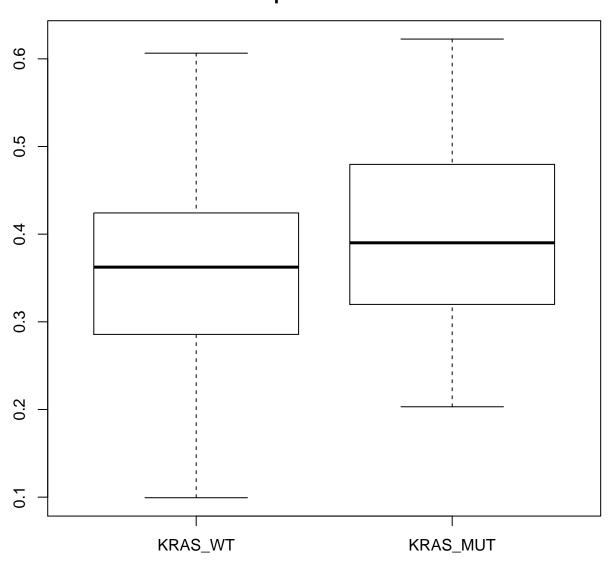
## krasqh\_200\_gene\_list/adap\_adap\_single p-val= 0.01



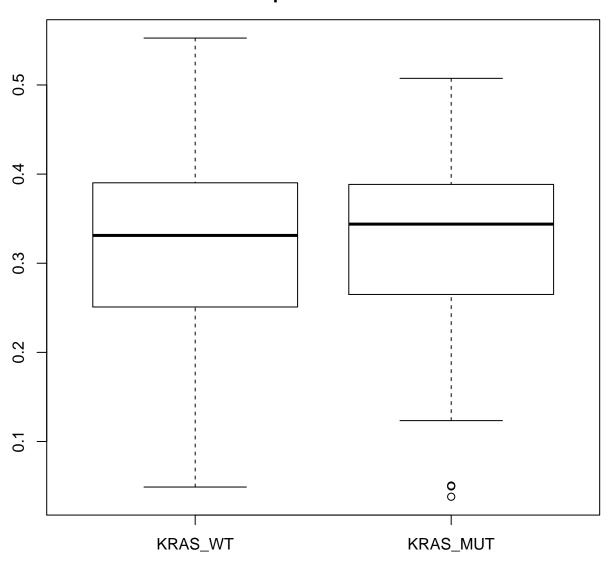
## krasqh\_300\_gene\_list/adapB\_single p-val= 0.03



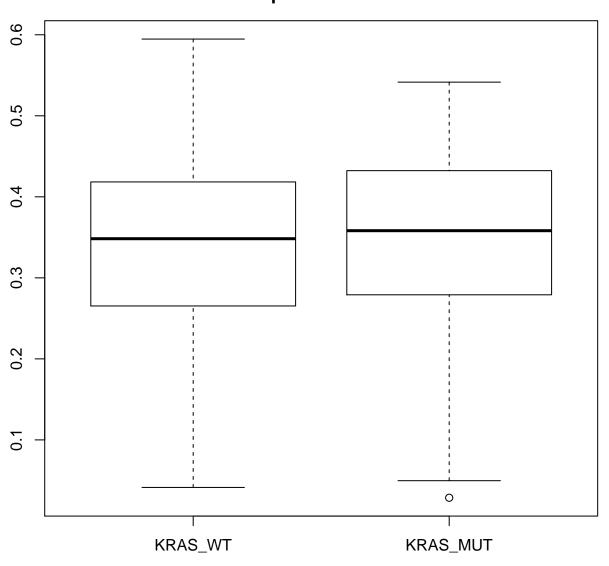
## krasqh\_300\_gene\_list/adap\_adap\_single p-val= 0.02



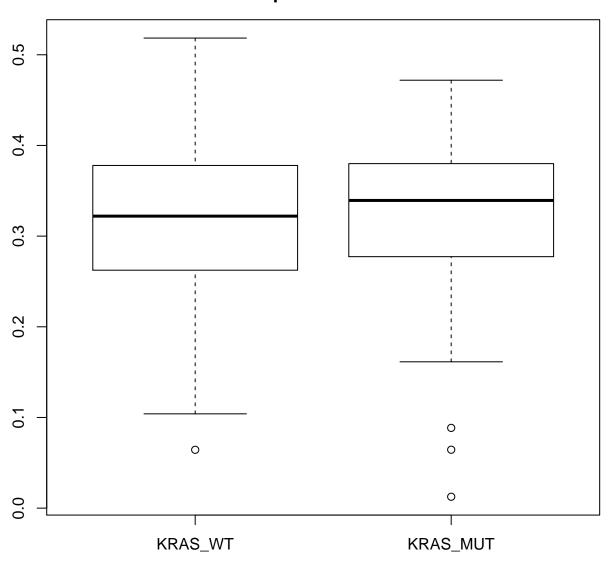
## kraswt\_100\_gene\_list/adapB\_single p-val= 0.92



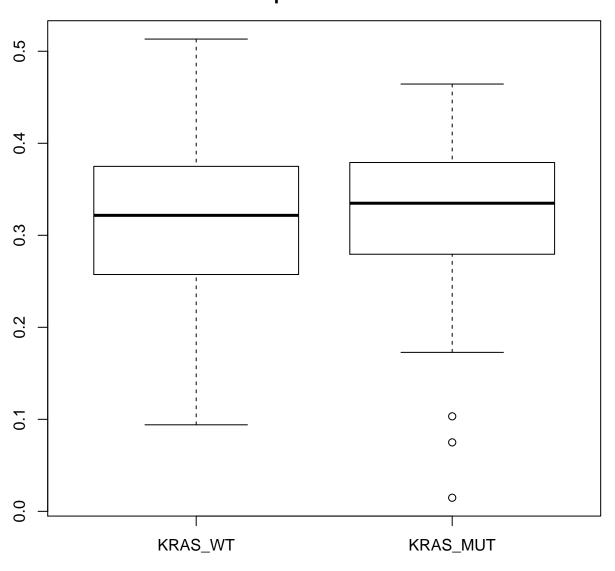
## kraswt\_100\_gene\_list/adap\_adap\_single p-val= 0.81



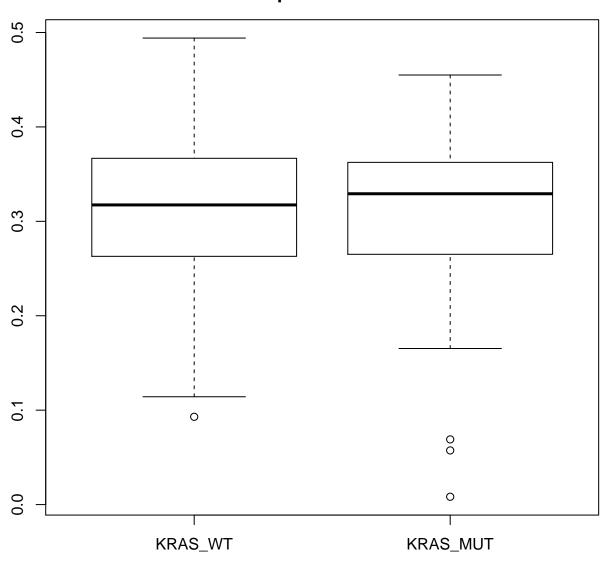
#### kraswt\_200\_gene\_list/adapB\_single p-val= 0.84



#### kraswt\_200\_gene\_list/adap\_adap\_single p-val= 0.76



## kraswt\_300\_gene\_list/adapB\_single p-val= 0.9



#### kraswt\_300\_gene\_list/adap\_adap\_single p-val= 0.98

