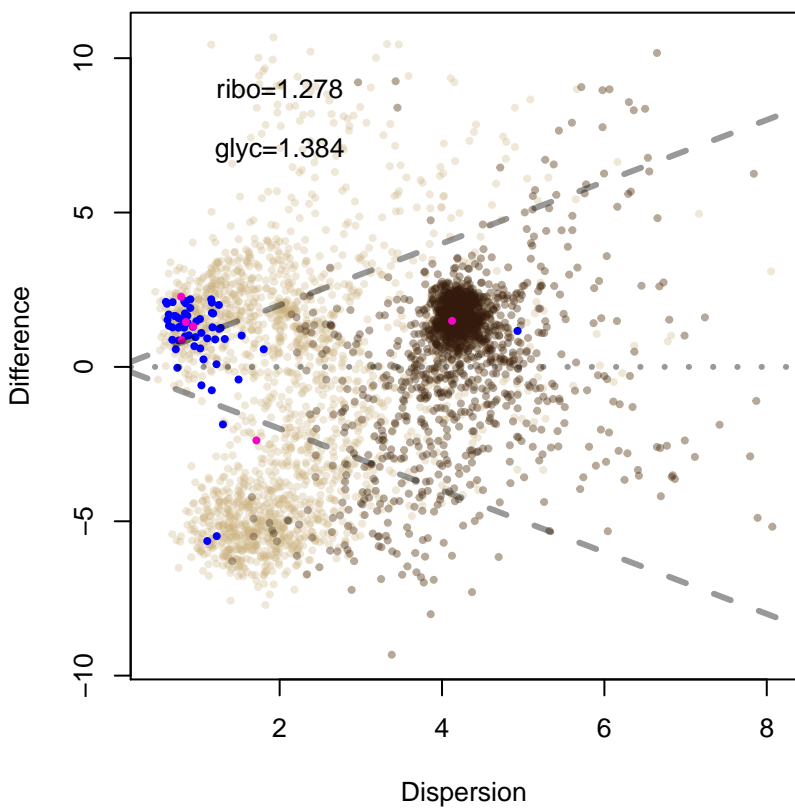
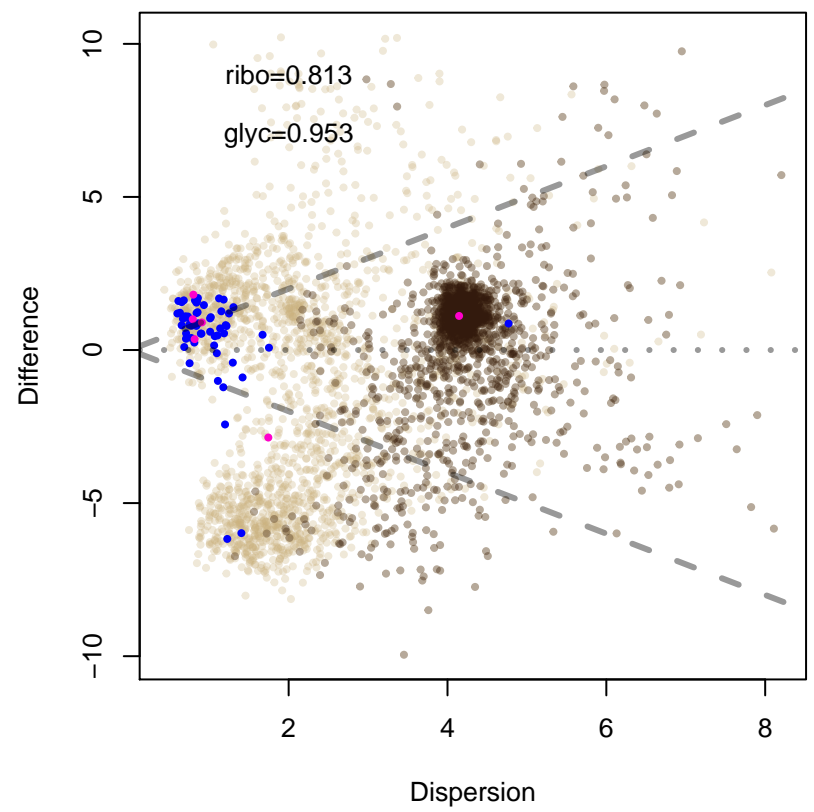


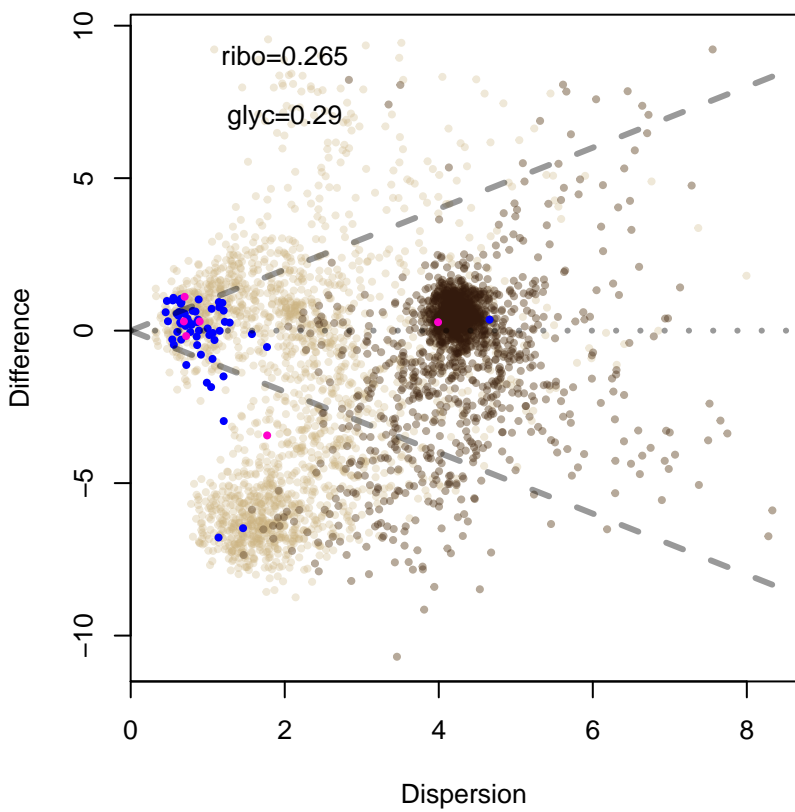
**denom = all**



**denom = IQLR**



**denom = LVHA**



**denom = ribosome**

