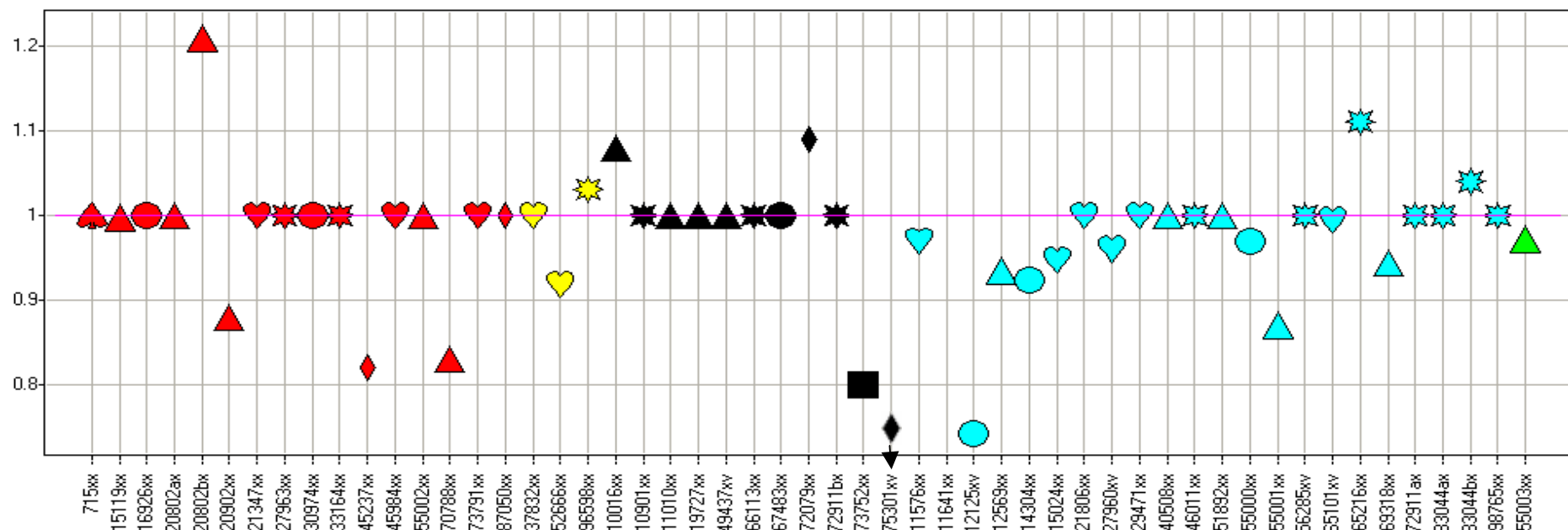
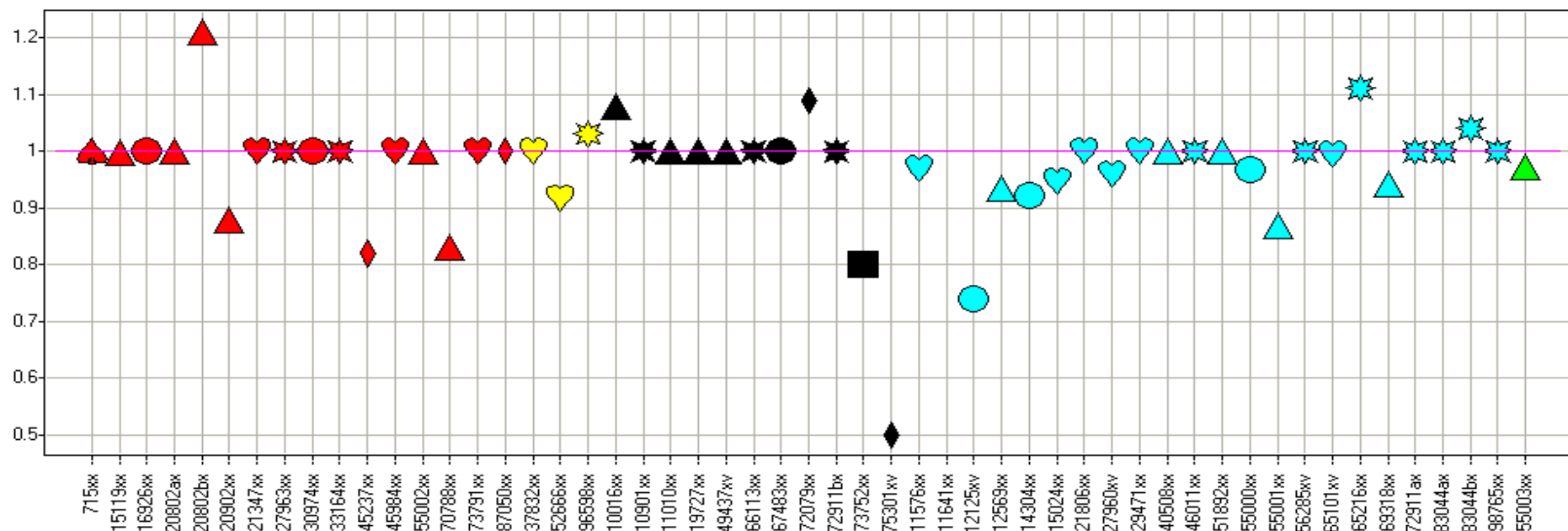
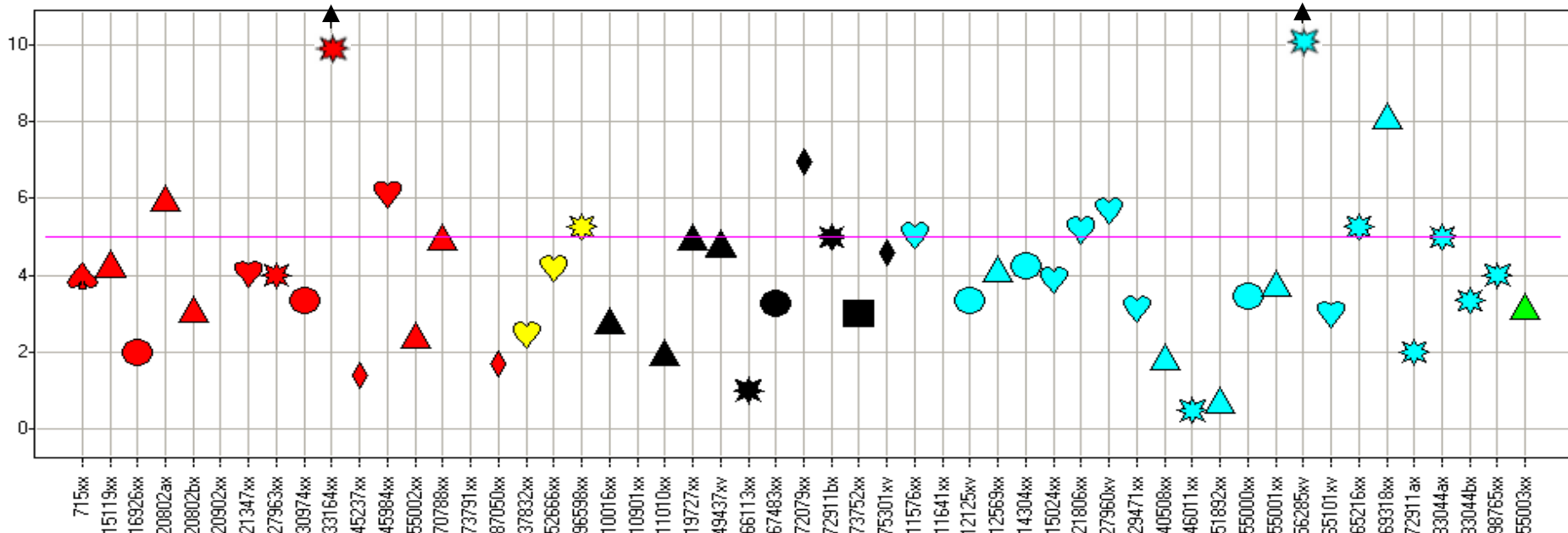
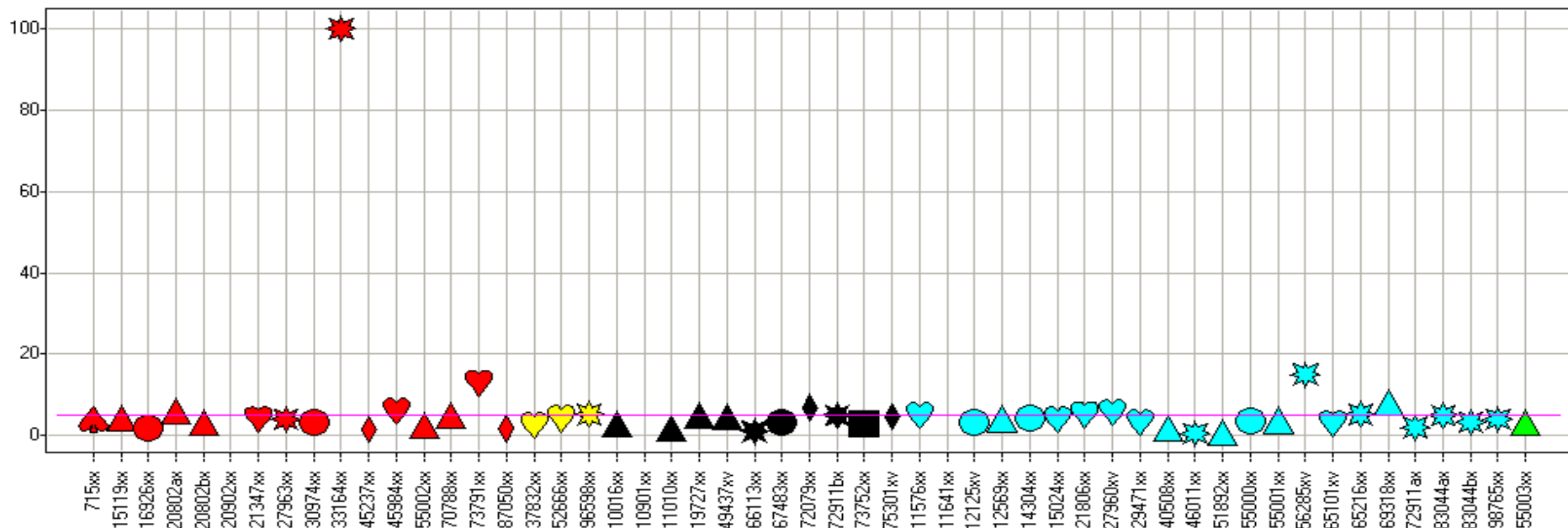


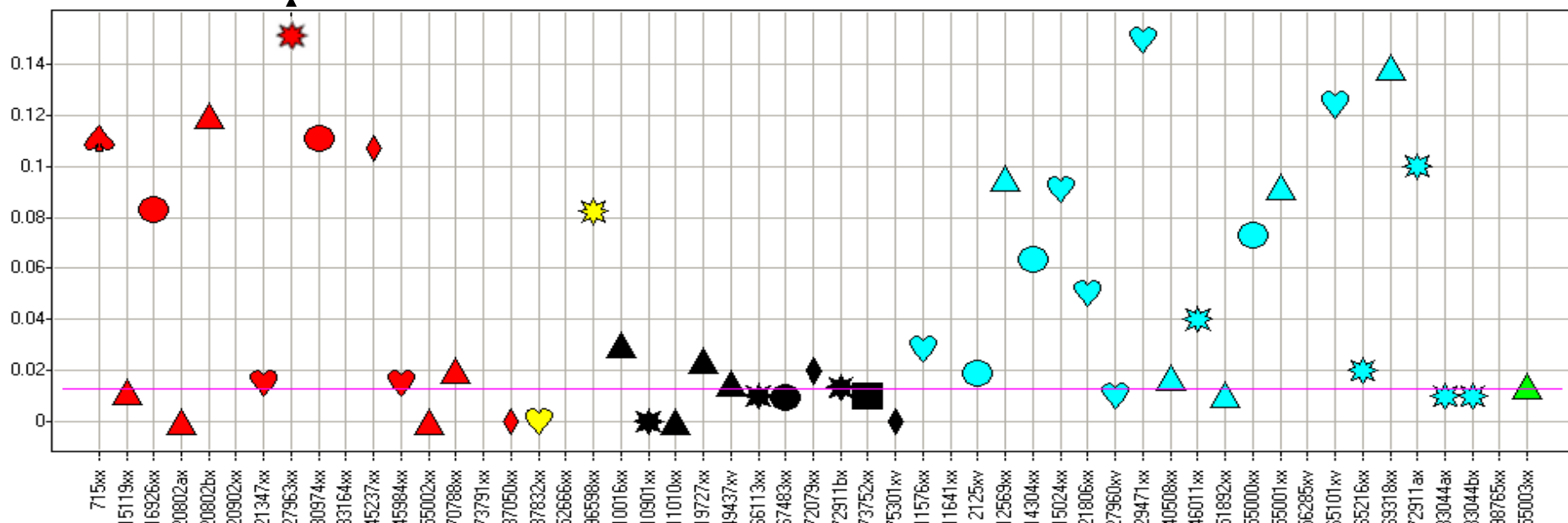
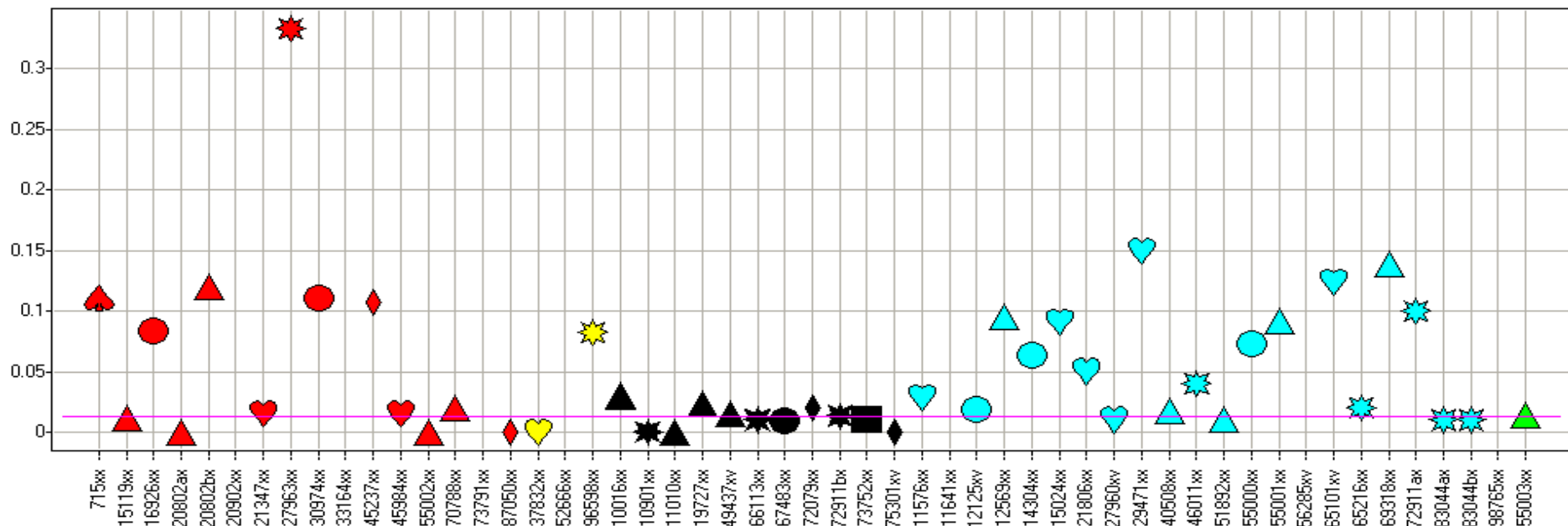
BSA (1:1)



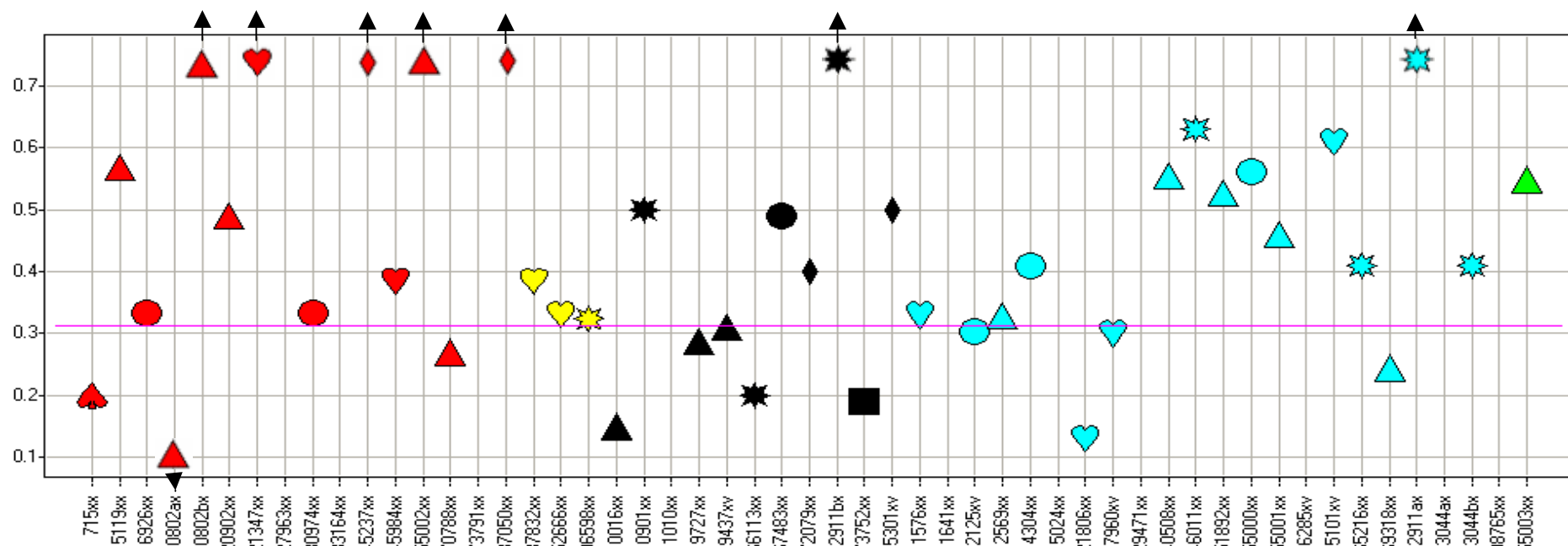
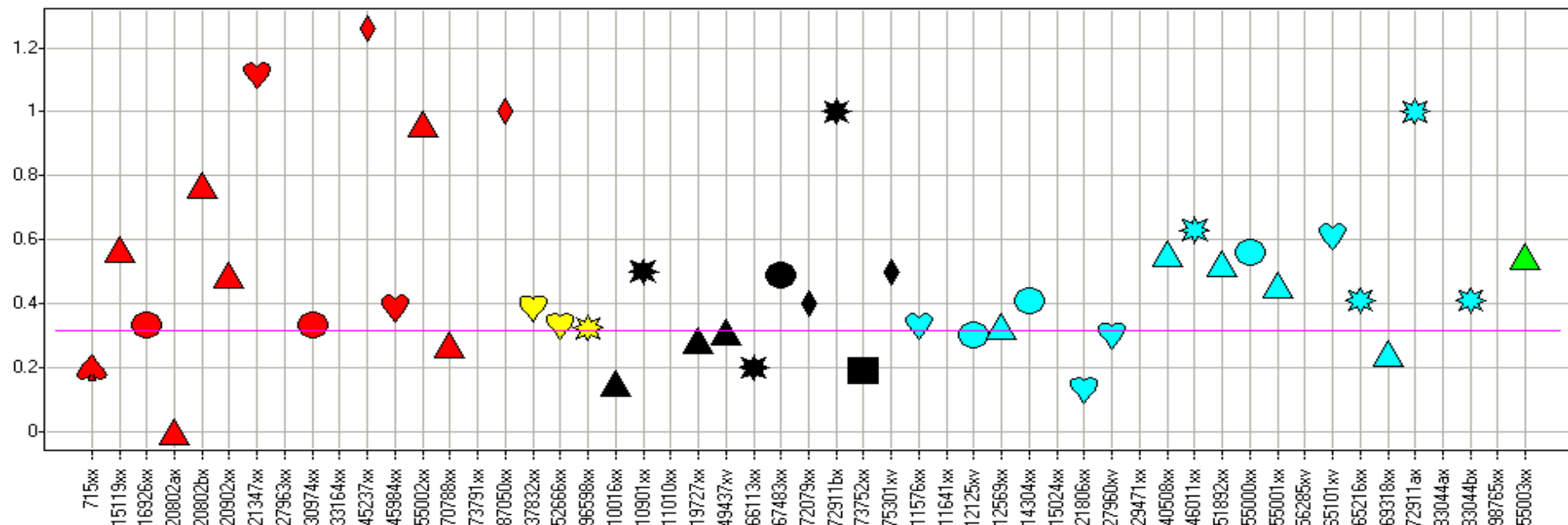
Catalase (5:1)



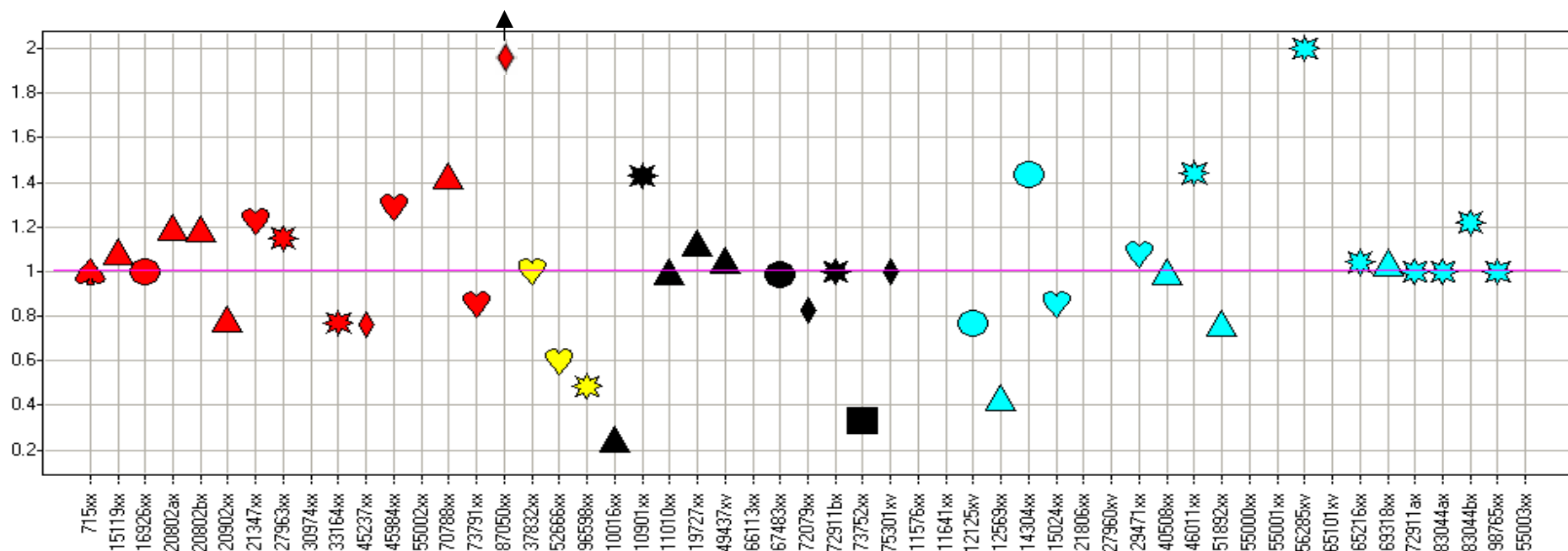
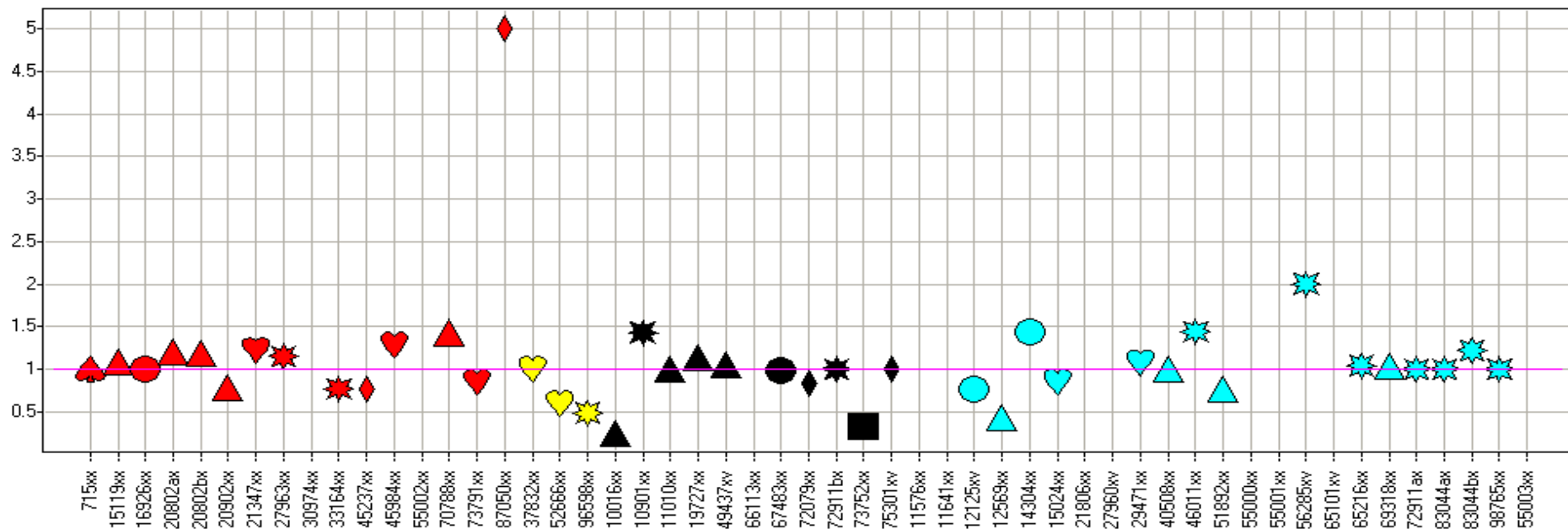
Glycogen phosphorylase (1:76)



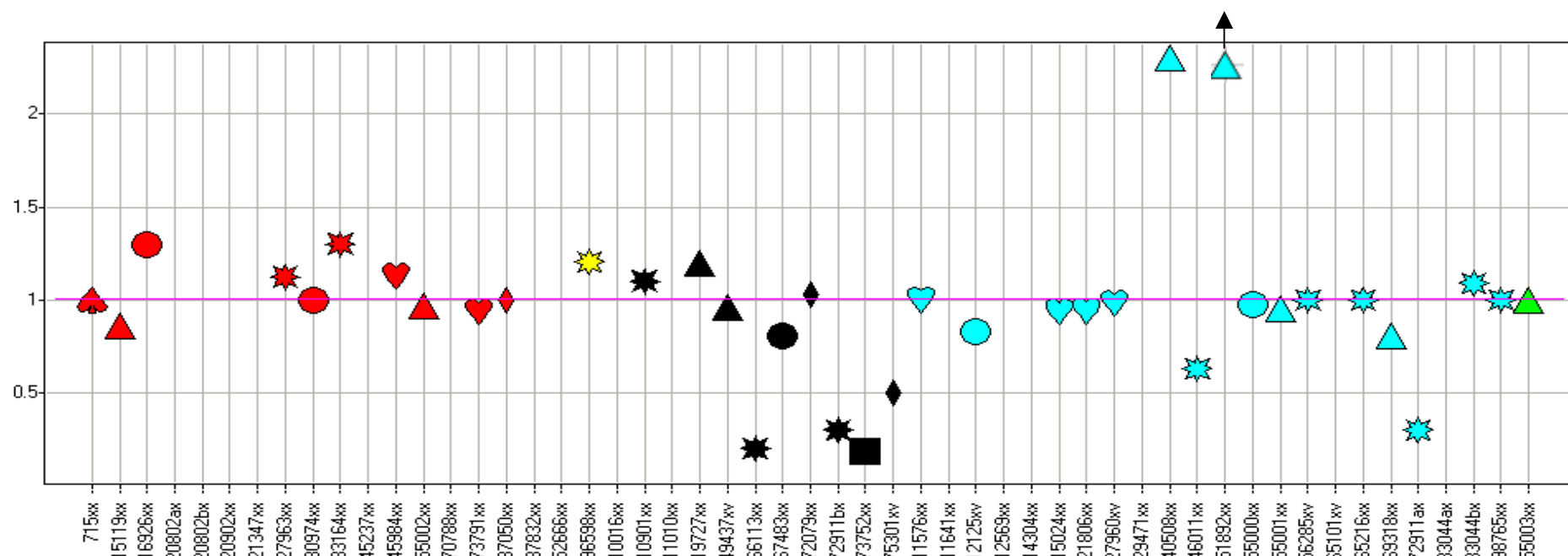
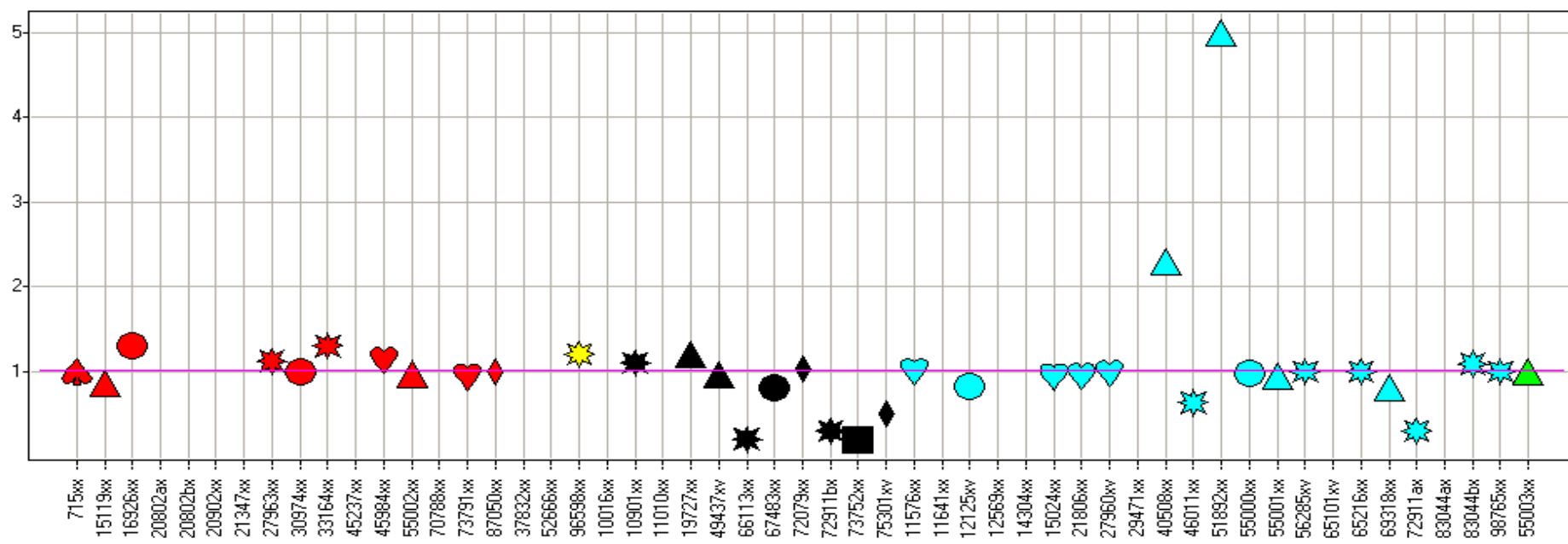
Carbonic Anhydrase (1:3)



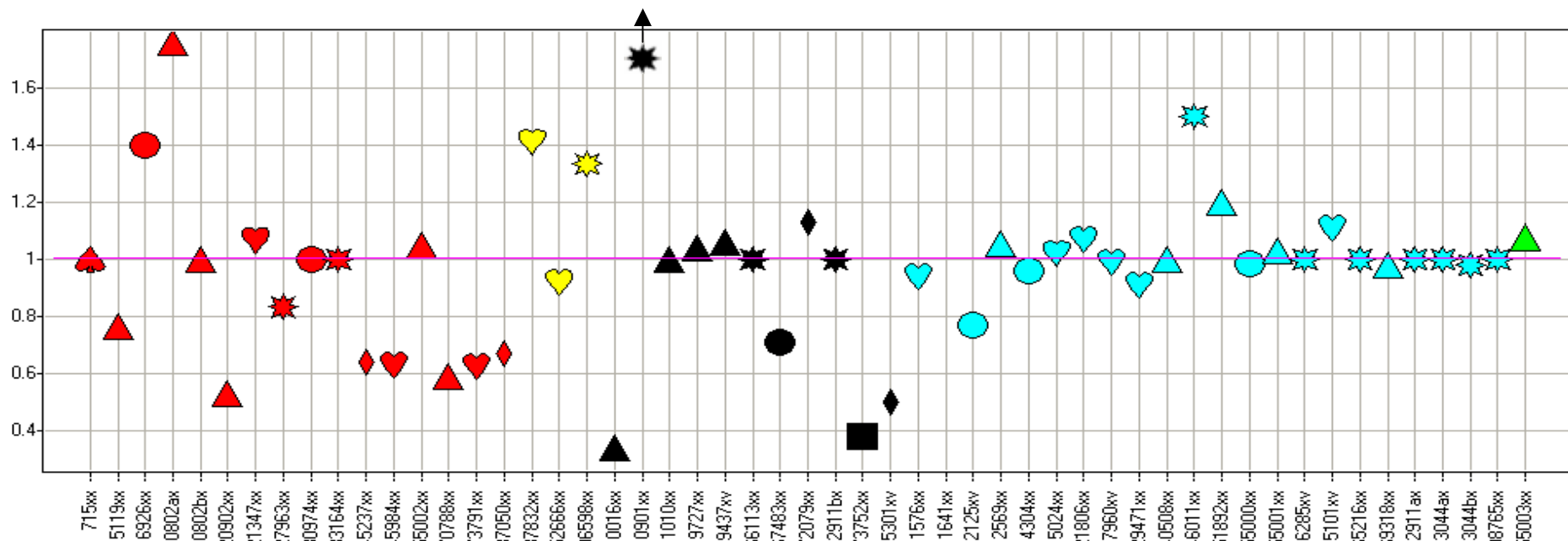
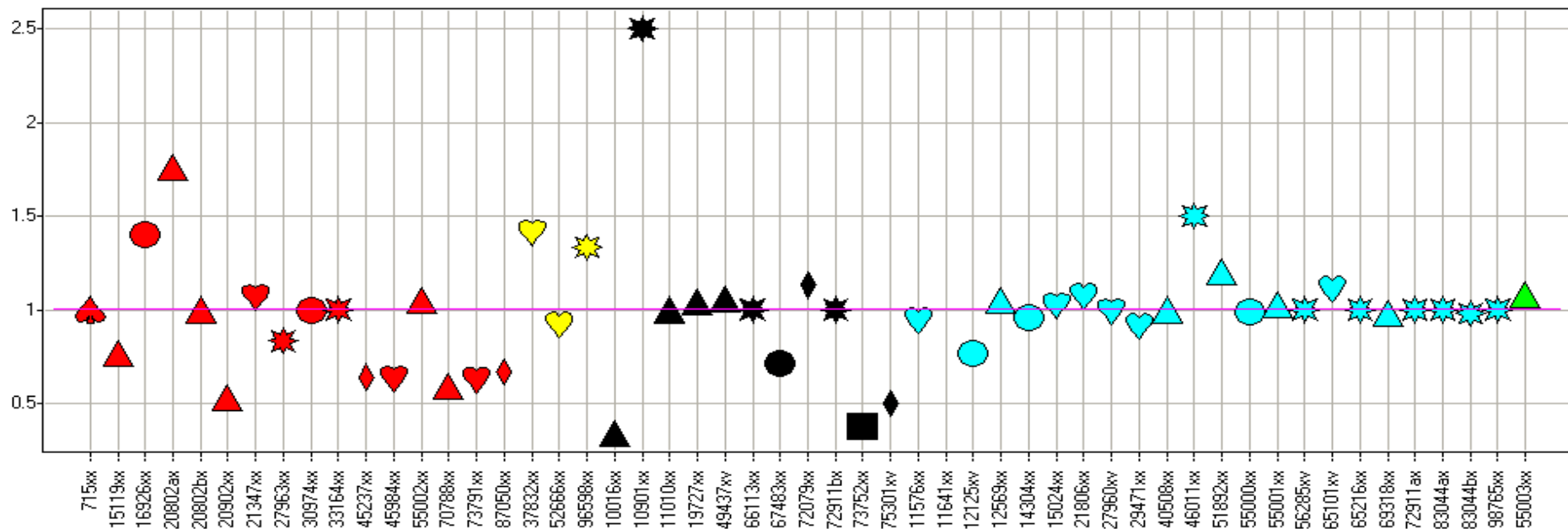
Peroxidase (1:1)



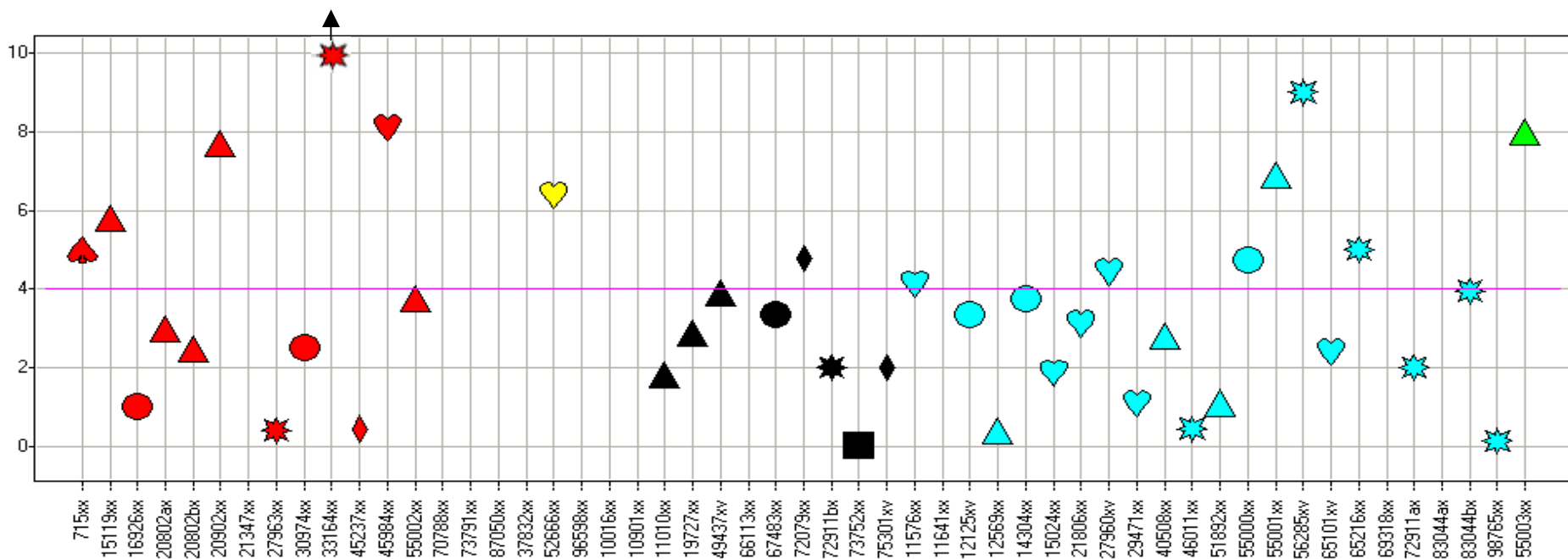
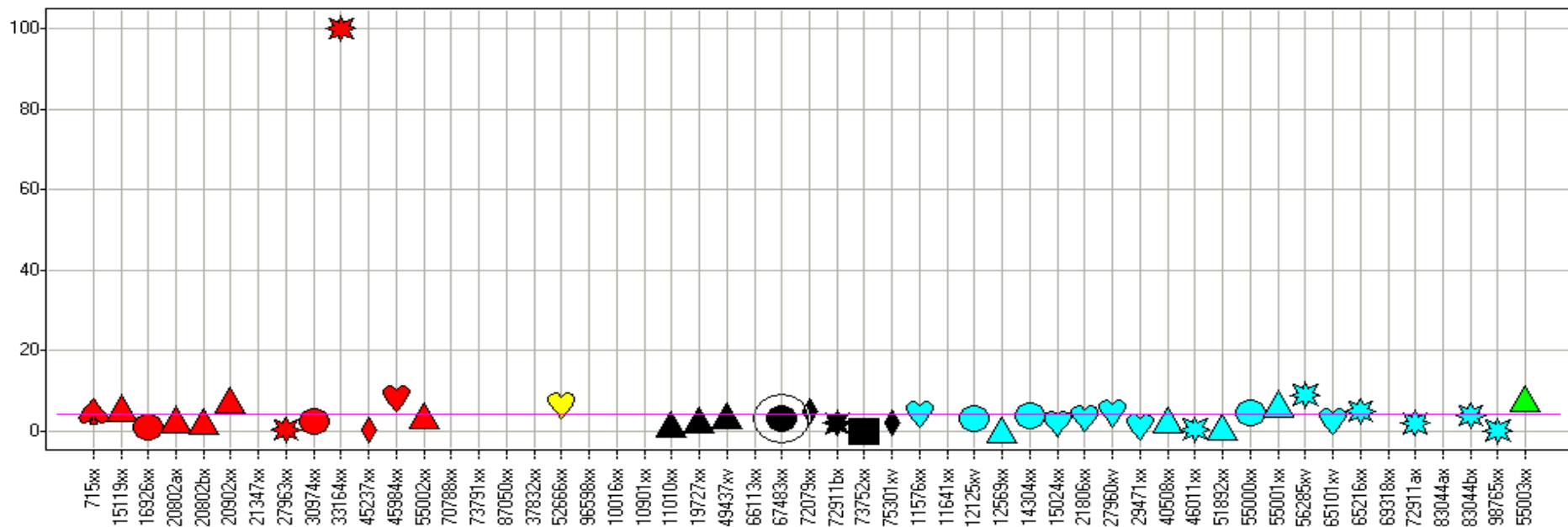
Ribonuclease A (1:1)



Lactoperoxidase (1:1)








beta Casein (4:1)






Legend

Method for Quantification

 Densitometry	 Ion current	 Spectral count
 DIGE	 Isotope MS	

Type of mass spectrometer

 3D ion trap	 Linear ion trap/FT	 TOF	 Combination
 Linear ion trap	 Quadrupole-TOF	 TOF/TOF	