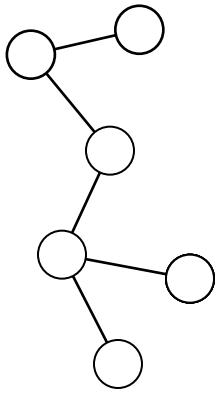
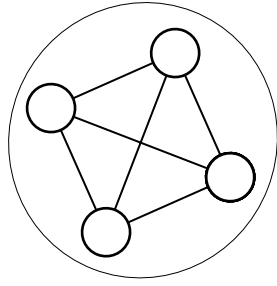


**Figure 1**

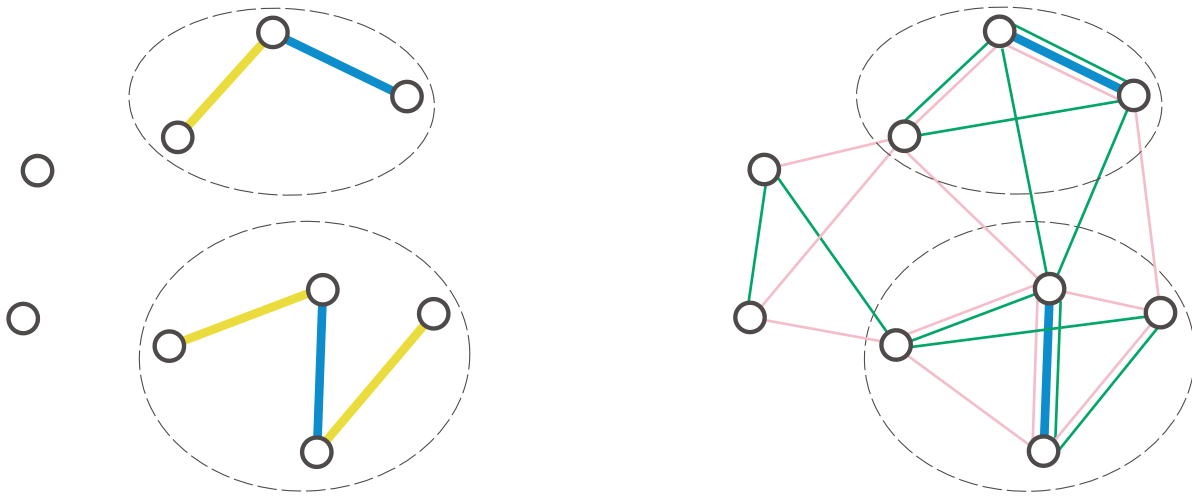


Binary interaction network

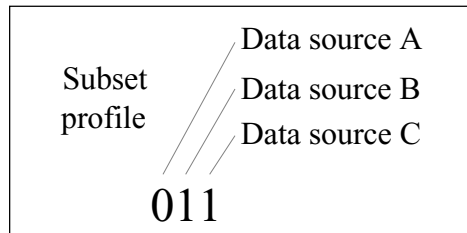
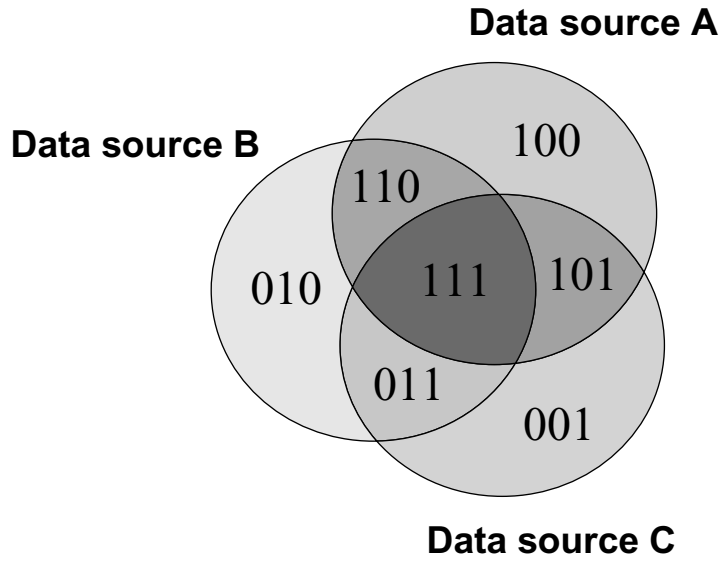


Protein complex with 4 subunits

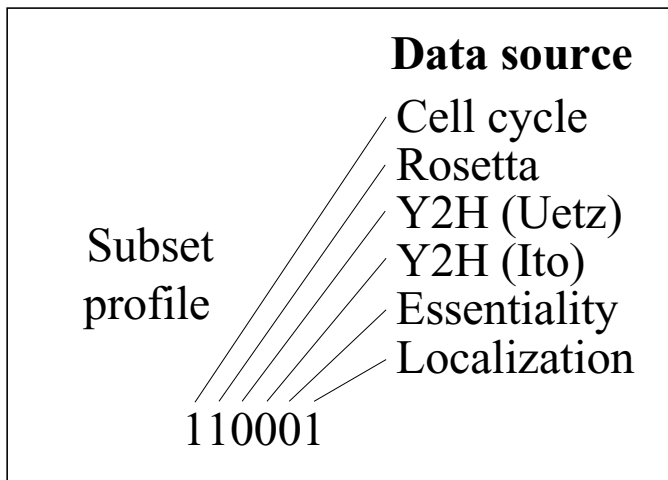
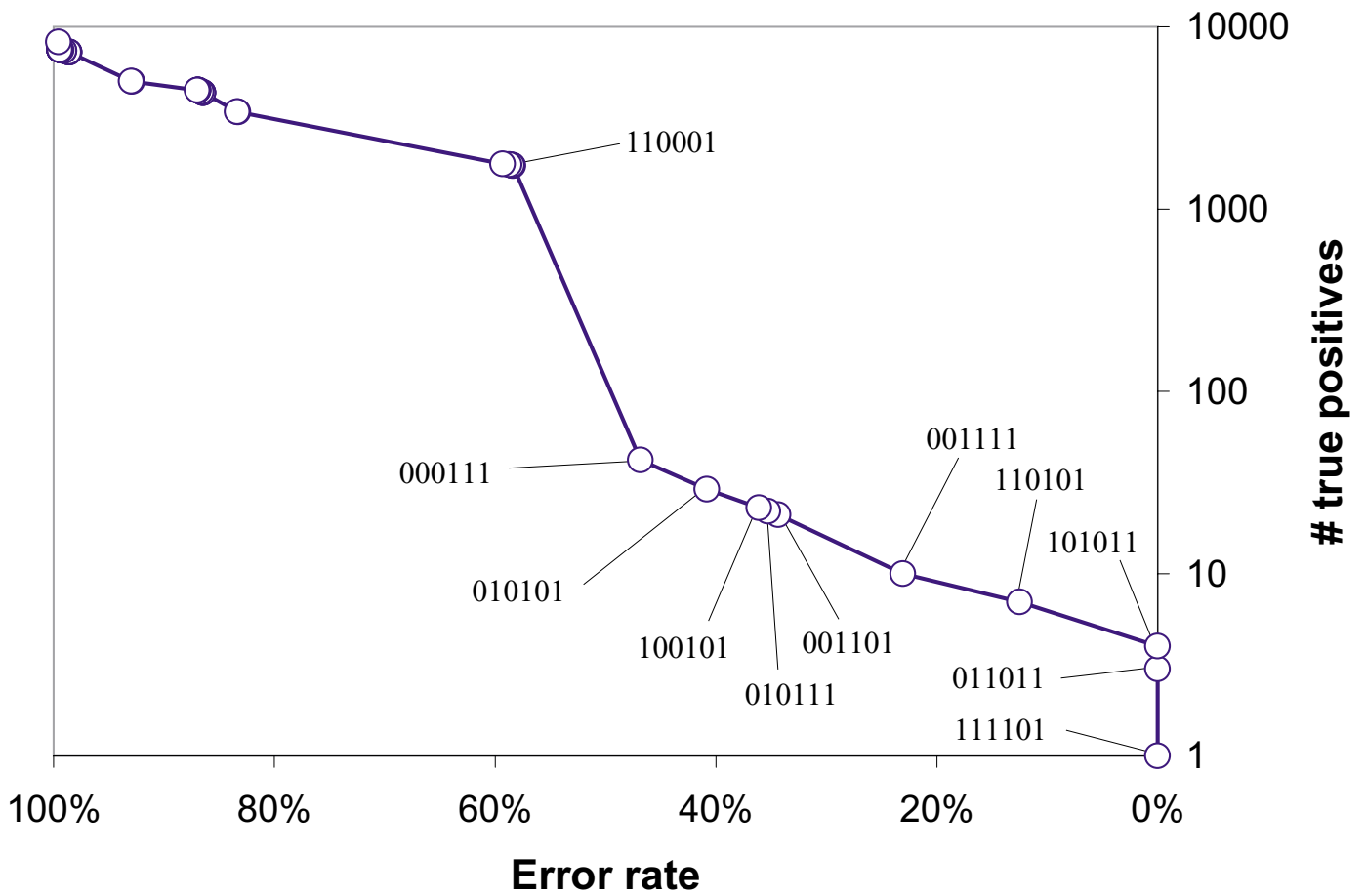
## Figure 2



**Figure 3**



**Figure 4**



**Figure 5**

<b>Data source</b>	<b>TP</b>	<b>FN</b>	<b>TN</b>	<b>FP</b>	<b>FN rate [%]</b>	<b>FP rate [%]</b>	<b>ε [%]</b>
Cell cycle	3,466	4,784	17,703,707	296,293	58.0%	1.6%	98.8
Rosetta	2,772	5,478	17,720,318	279,682	66.4%	1.6%	99.0
Y2H (Uetz)	50	8,200	17,999,109	891	99.4%	0.0%	94.7
Y2H (Ito)	78	8,172	17,995,806	4,194	99.1%	0.0%	98.2
Essentiality	2,718	5,532	15,520,460	2,479,540	67.1%	13.8%	99.9
Localization	7,220	1,030	17,523,304	476,696	12.5%	2.6%	98.5

**Table 1**