

	Common Clusters (1)		Common Clusters (2)	
Trithorax-ASH2	6	75%	4	50%
Trithorax-NURF	4	57%	2	29%
Trithorax-dMyc	6	100%	6	100%
Trithorax-ASH1	4	50%	2	25%
		71%		57%

Common clusters (1): cluster overlap considering genes affected in the same or the opposite way in the mutants analyzed. Common clusters (2): cluster overlap only considering genes affected in the same way in the mutants analyzed.