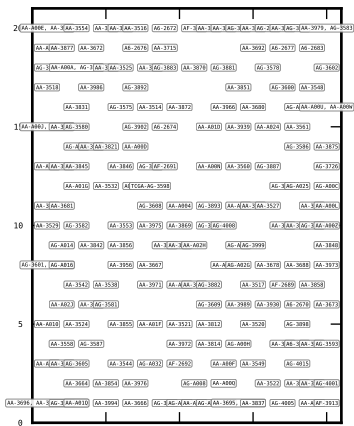
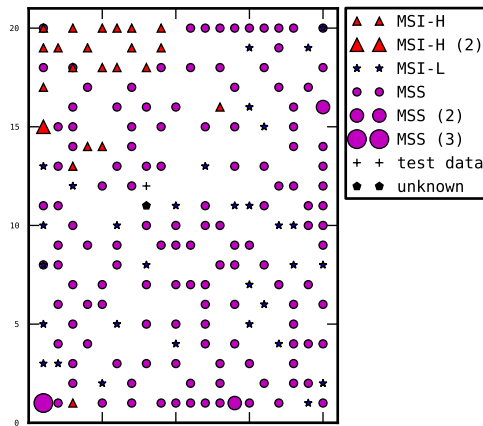


number of training samples	: 193	map rows	: 20
number of test samples	: 1	map cols	: 20
features size	: 17719	max neighborhood size	: 15
training iterations	: 50	neighborhood step size	: 0.3
		random seed	: 444807381

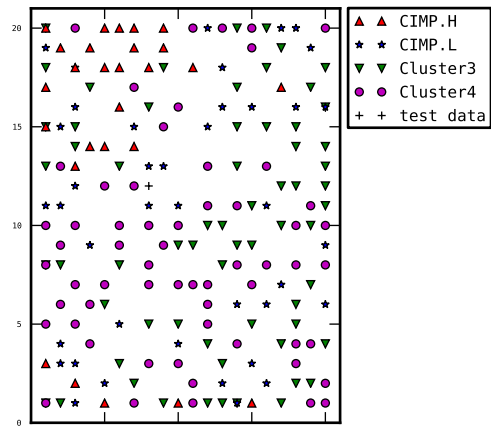
samples name



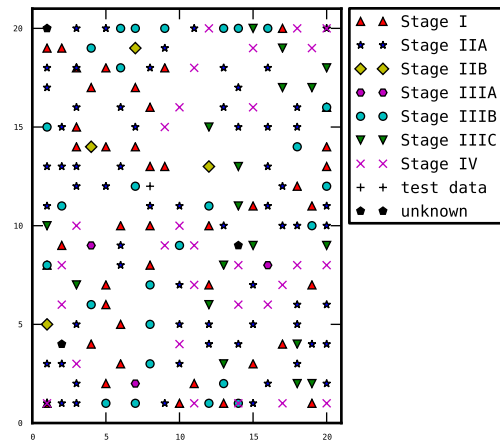
MSI_status



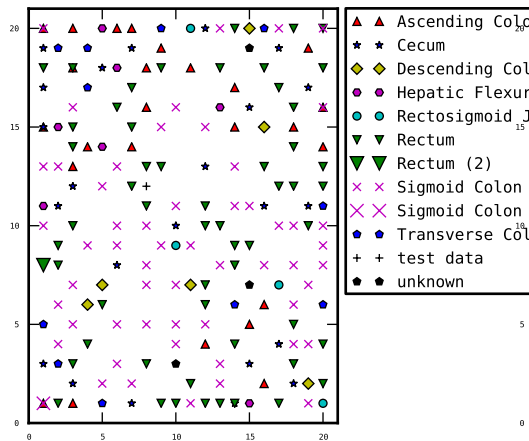
methylation_subtype



tumor_stage



anatomic_organ_subdivision



tumor_site

