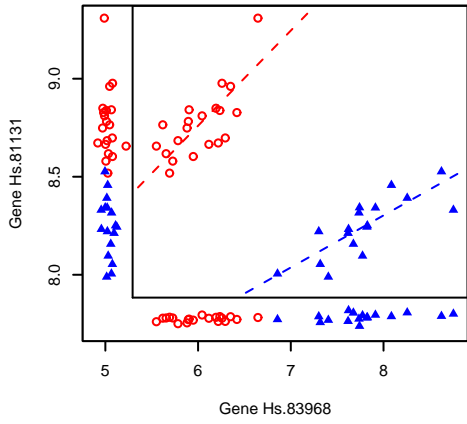
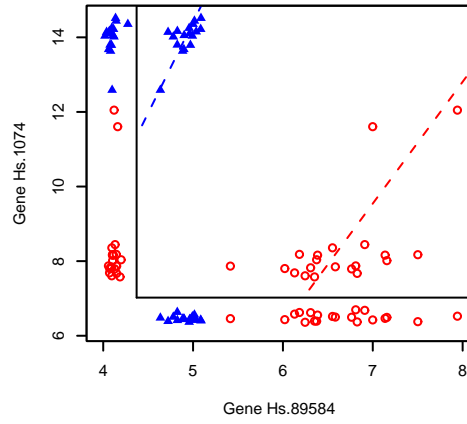


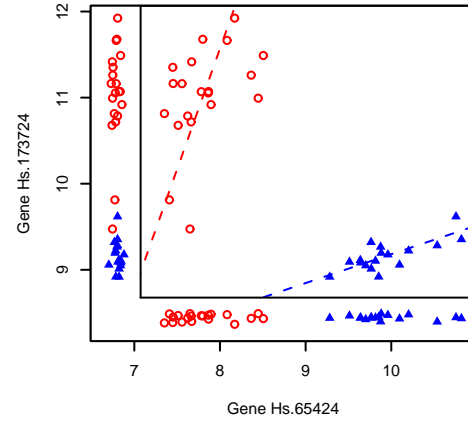
Gene Pair 1 , Score = 2.46



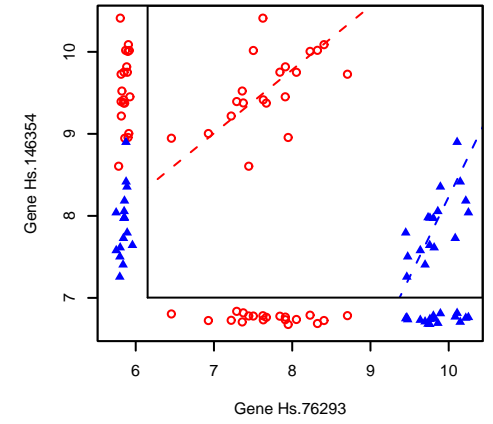
Gene Pair 2 , Score = 2.43



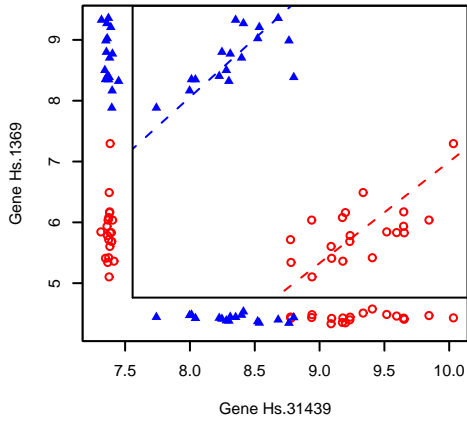
Gene Pair 3 , Score = 2.39



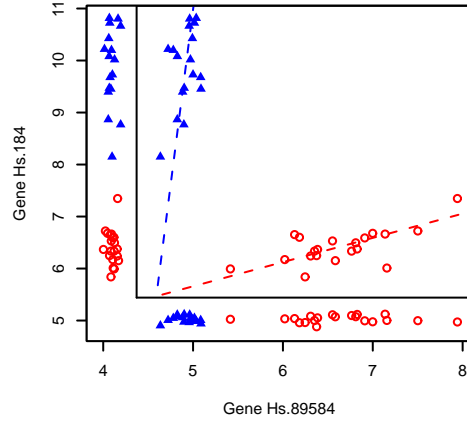
Gene Pair 4 , Score = 2.34



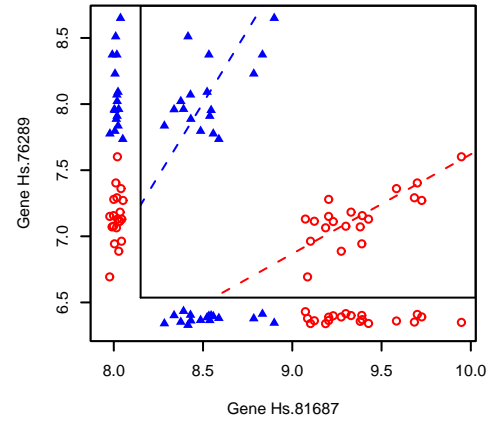
Gene Pair 5 , Score = 2.34



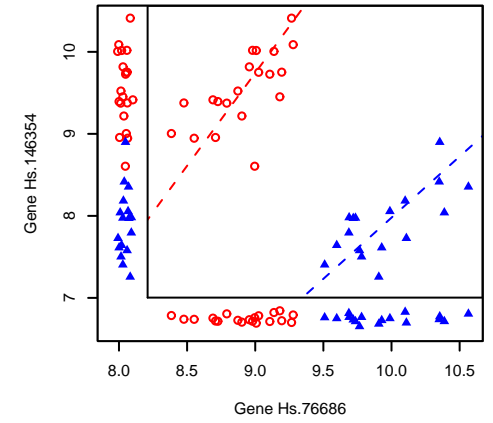
Gene Pair 6 , Score = 2.31



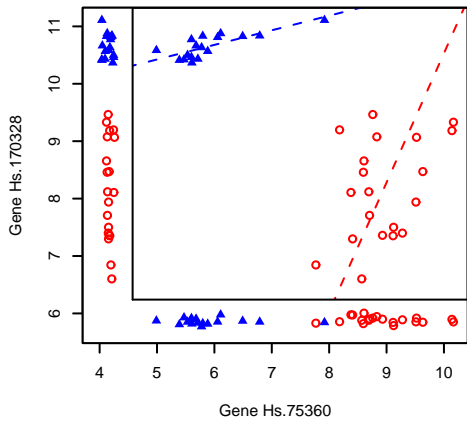
Gene Pair 7 , Score = 2.31



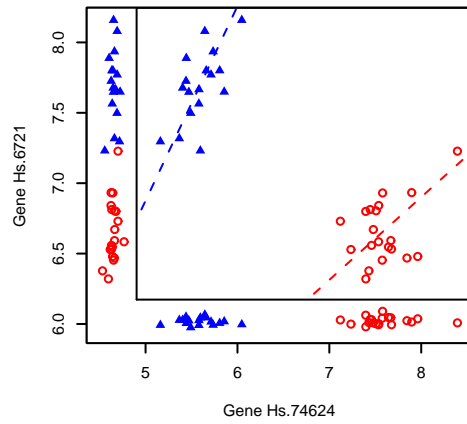
Gene Pair 8 , Score = 2.3



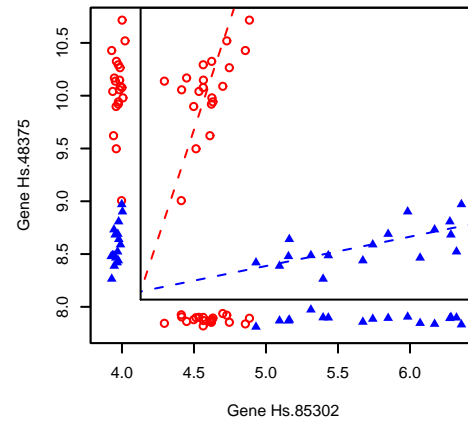
Gene Pair 9 , Score = 2.3



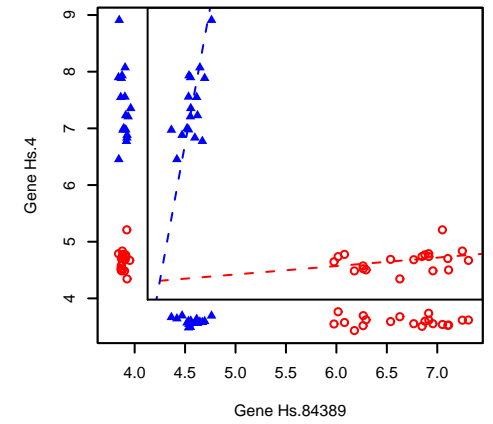
Gene Pair 10 , Score = 2.28



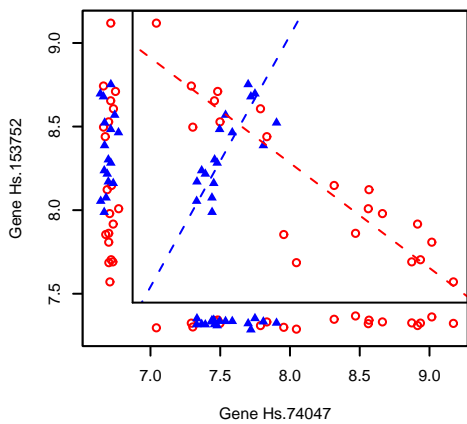
Gene Pair 11 , Score = 2.28



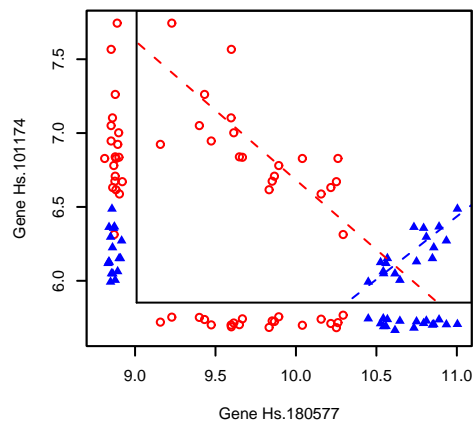
Gene Pair 12 , Score = 2.27



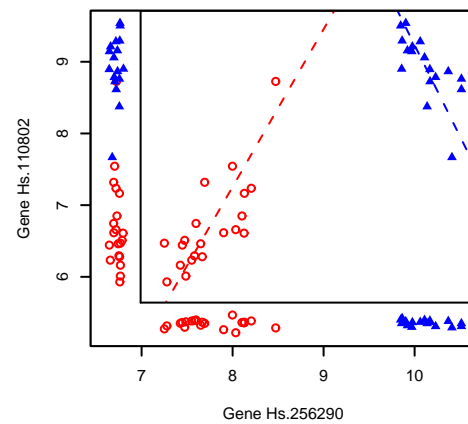
Gene Pair 1 , Score = 1.65



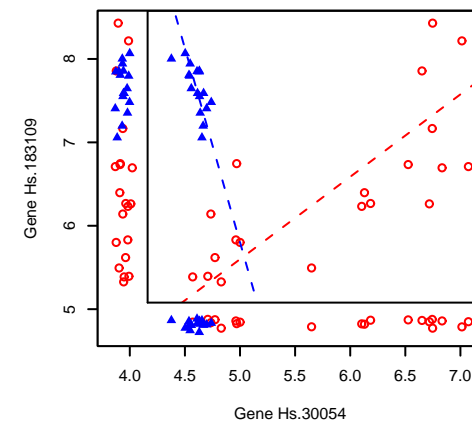
Gene Pair 2 , Score = 1.59



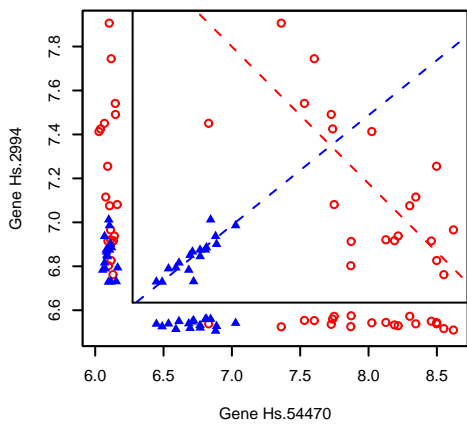
Gene Pair 3 , Score = 1.59



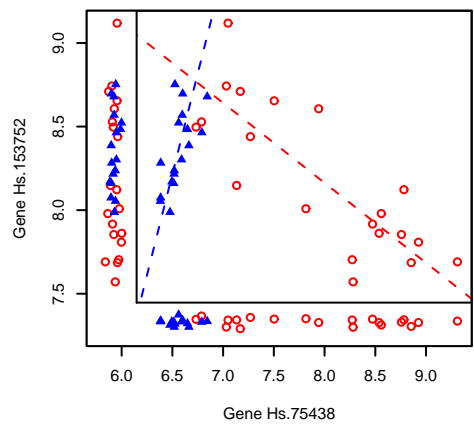
Gene Pair 4 , Score = 1.57



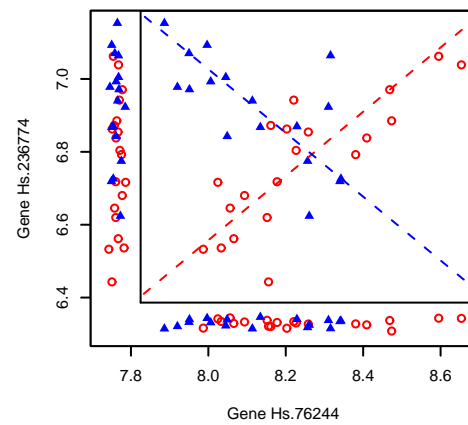
Gene Pair 5 , Score = 1.57



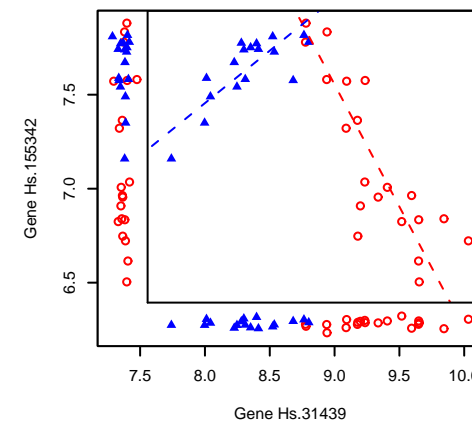
Gene Pair 6 , Score = 1.54



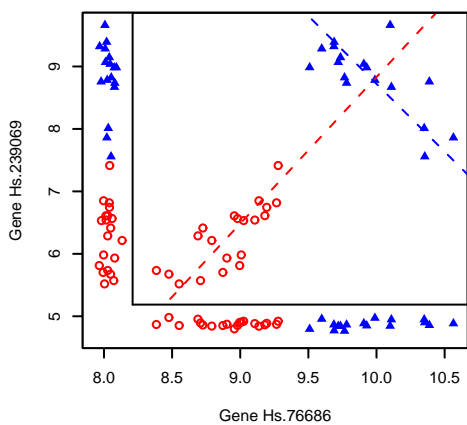
Gene Pair 7 , Score = 1.54



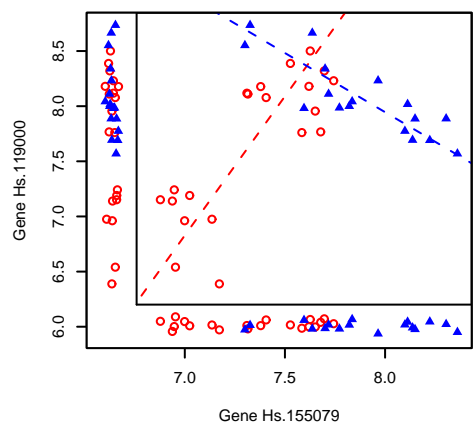
Gene Pair 8 , Score = 1.53



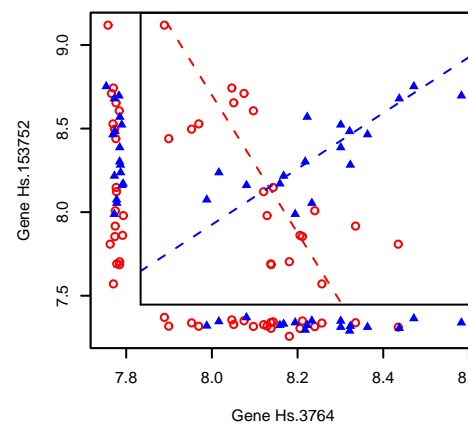
Gene Pair 9 , Score = 1.53



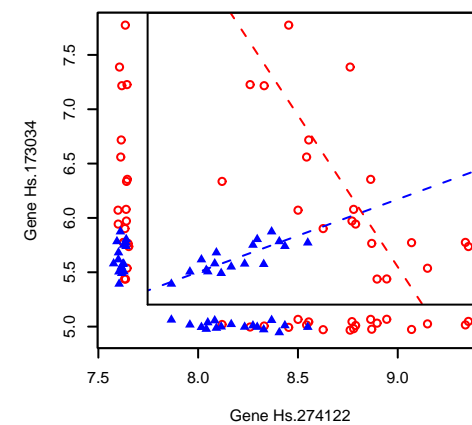
Gene Pair 10 , Score = 1.53



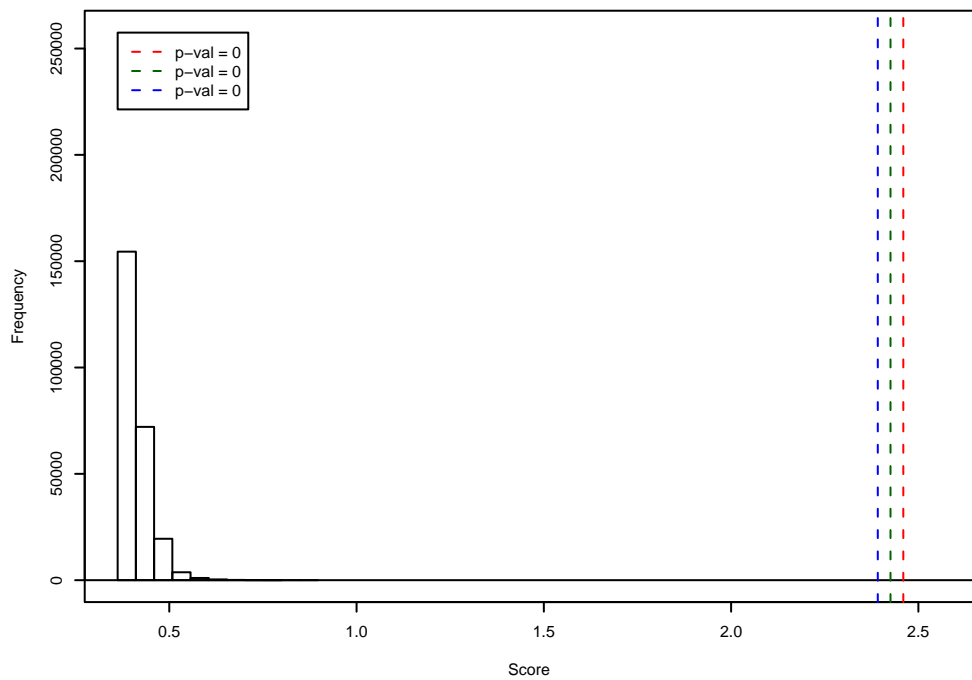
Gene Pair 11 , Score = 1.53



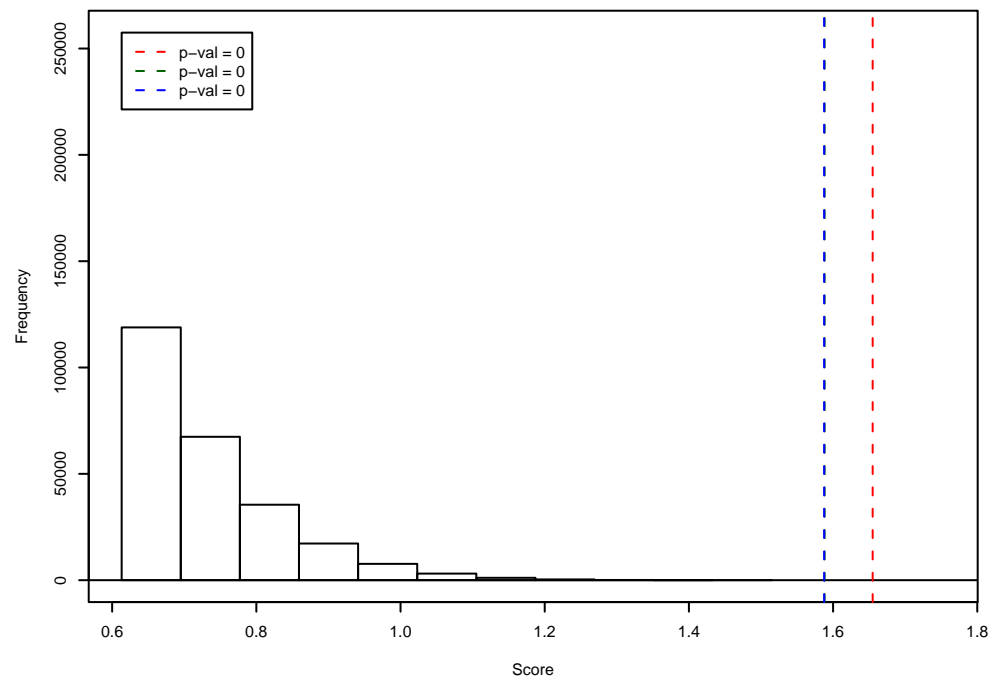
Gene Pair 12 , Score = 1.52



Gap/Substitution



On/Off



p-value	# gp exceeding	% gp exceeding	fold advantage
0e+00	351008	6.984	Inf
1e-06	498813	9.925	99245.83
1e-05	674817	13.426	13426.43
1e-04	922033	18.345	1834.51
1e-03	1243663	24.744	247.44

p-value	# gp exceeding	% gp exceeding	fold advantage
0e+00	14	0.000	Inf
1e-06	81	0.002	16.12
1e-05	481	0.010	9.57
1e-04	3045	0.061	6.06
1e-03	18805	0.374	3.74