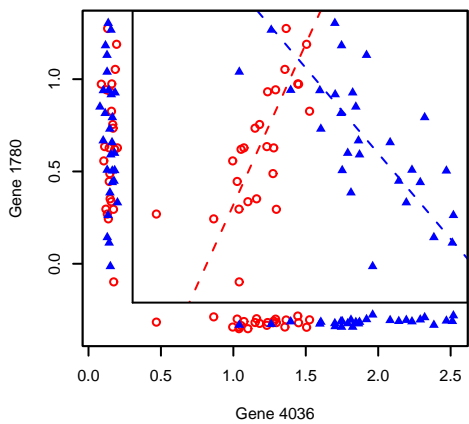
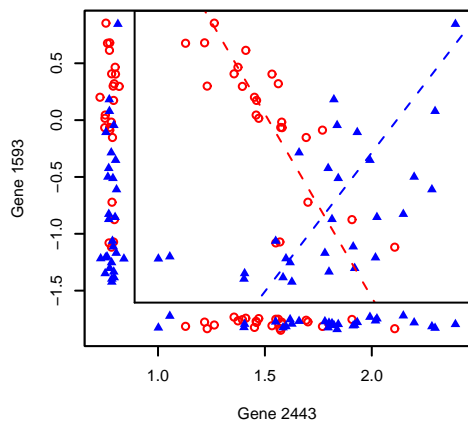


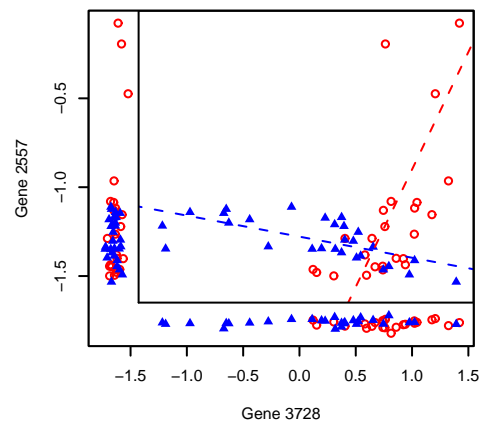
Gene Pair 1 , Score = 1.5



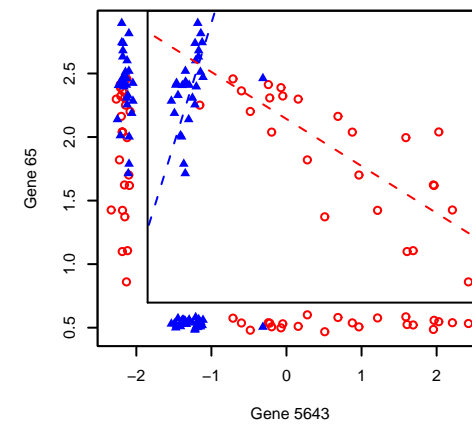
Gene Pair 2 , Score = 1.48



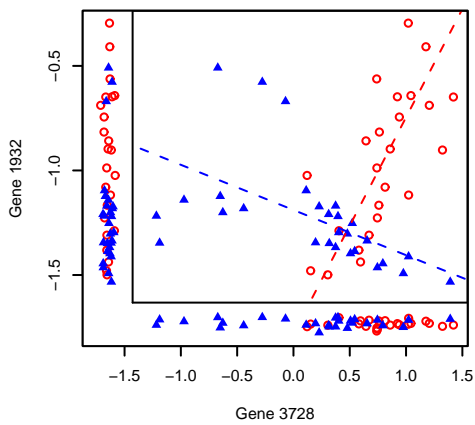
Gene Pair 3 , Score = 1.45



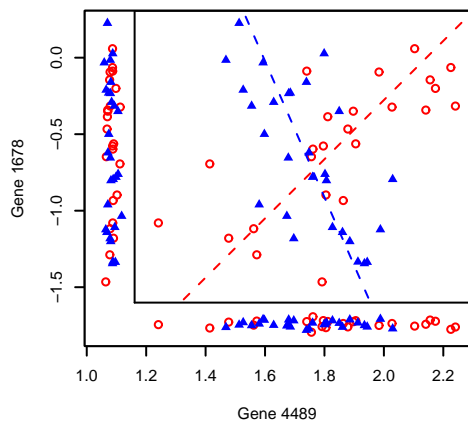
Gene Pair 4 , Score = 1.44



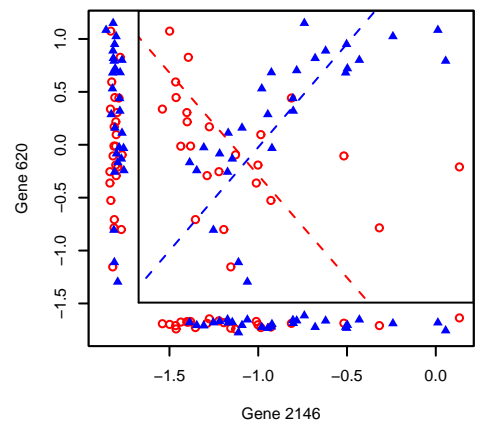
Gene Pair 5 , Score = 1.44



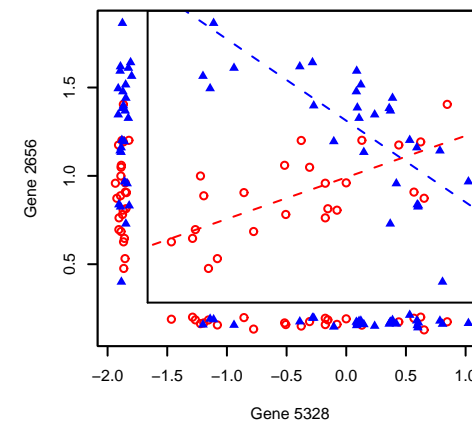
Gene Pair 6 , Score = 1.43



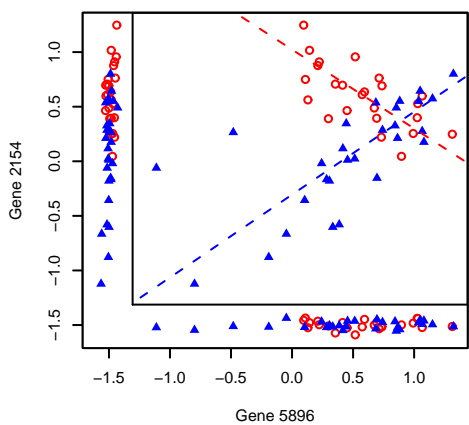
Gene Pair 7 , Score = 1.43



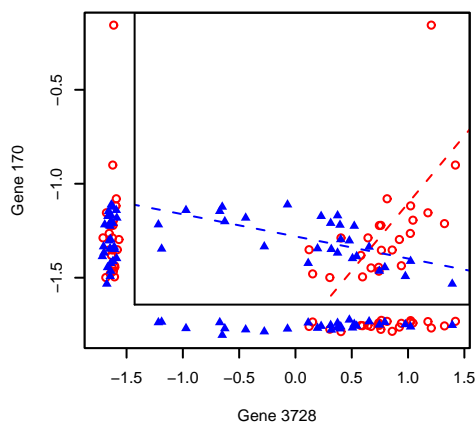
Gene Pair 8 , Score = 1.42



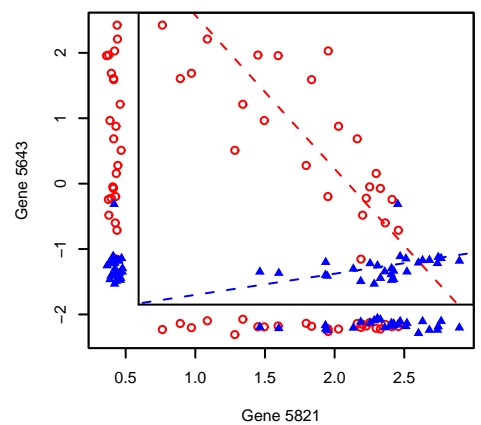
Gene Pair 9 , Score = 1.42



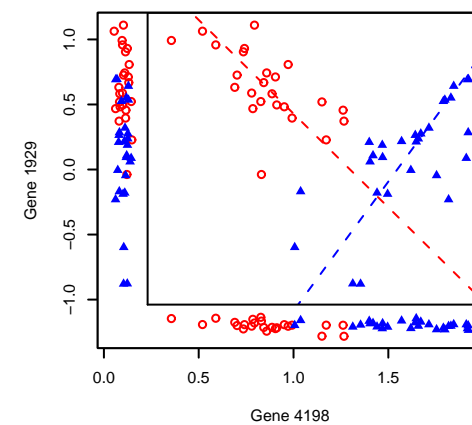
Gene Pair 10 , Score = 1.41



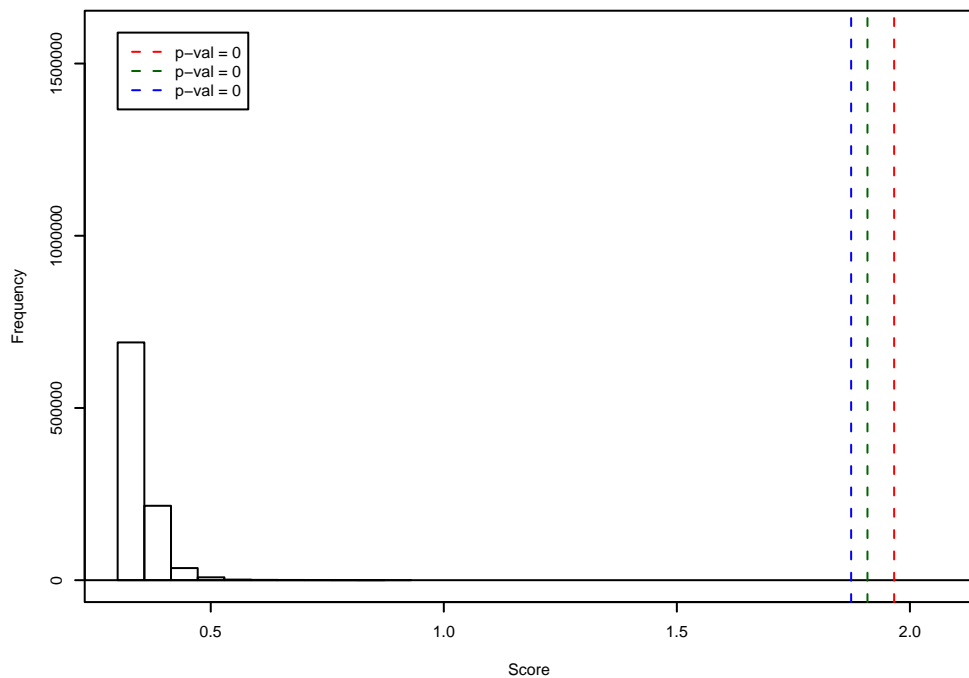
Gene Pair 11 , Score = 1.4



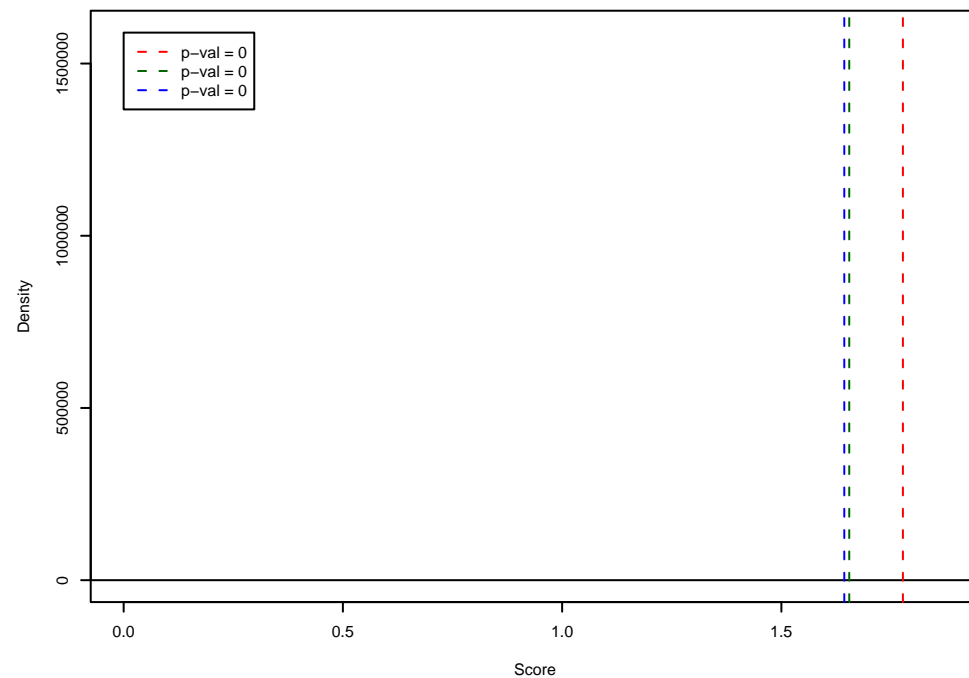
Gene Pair 12 , Score = 1.4



Gap/Substitution



On/Off



p-value	# gp exceeding	% gp exceeding	fold advantage
0e+00	138841	0.728	Inf
1e-06	209154	1.097	10965.07
1e-05	346638	1.817	1817.28
1e-04	1193706	6.258	625.81
1e-03	2889226	15.147	151.47

p-value	# gp exceeding	% gp exceeding	fold advantage
0e+00	323	0.002	Inf
1e-06	1997	0.010	104.69
1e-05	6640	0.035	34.81
1e-04	25284	0.133	13.26
1e-03	116386	0.610	6.10