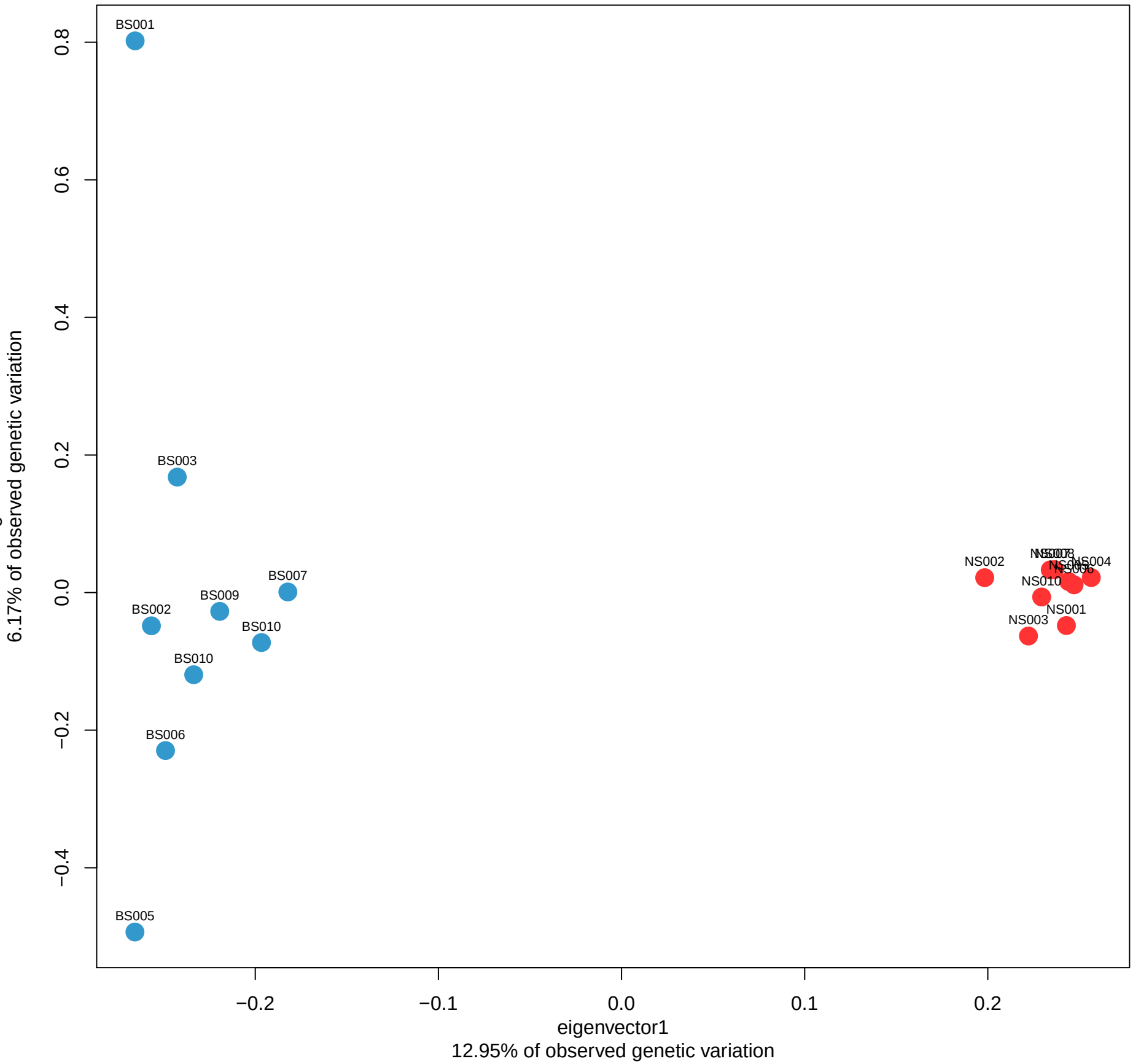
BS001
BS003
BS007
BS009
BS002
BS010
BS010
BS006
BS005
NS002
NS004
NS010
NS001
NS003
0.8
0.6
0.4
0.2
0.0
−0.2
−0.4
6.17% of observed genetic variation
−0.2
−0.1
0.0
0.1
0.2
eigenvector1
12.95% of observed genetic variation