



clust\_2374: Type I restriction–modification system, specificity subunit S (EC 3.1.21.3)

clust\_2255: "Type I restriction–modification system, specificity subunit S "

clust\_2073: Type III restriction–modification system methylation subunit (EC 2.1.1.72)

clust\_1902: Type I restriction–modification system, specificity subunit S (EC 3.1.21.3)

clust\_1804: FIG045374: Type II restriction enzyme, methylase subunit YeeA

clust\_1801: Type II restriction enzyme DpnII (EC 3.1.21.4) (Endonuclease DpnII) (R.DpnII)

clust\_1748: Type I restriction–modification system, specificity subunit S (EC 3.1.21.3)

clust\_1629: Type II restriction endonuclease

clust\_1612: Type III restriction–modification system methylation subunit (EC 2.1.1.72)

clust\_1464: Type I restriction–modification system, restriction subunit R (EC 3.1.21.3)

clust\_1439: Type I restriction–modification system, specificity subunit S (EC 3.1.21.3)

clust\_1430: Type III restriction–modification system DNA endonuclease res (EC 3.1.21.5)

clust\_1397: "Type I restriction–modification system, specificity subunit S "

clust\_1366: FIG045374: Type II restriction enzyme, methylase subunit YeeA

clust\_1321: Type II restriction endonuclease

clust\_1290: Type I restriction–modification system, DNA–methyltransferase subunit M (EC 2.1.1.72)

clust\_1163: Type I restriction–modification system, specificity subunit S (EC 3.1.21.3)

clust\_1162: Type I restriction–modification system, DNA–methyltransferase subunit M (EC 2.1.1.72)

clust\_1105: Type I restriction–modification system, restriction subunit R (EC 3.1.21.3)

clust\_1089: Type I restriction–modification system, restriction subunit R (EC 3.1.21.3)