Custom edge-adjusted network - all nodes

2ARC.A_@aC_E.coli



Custom edge-adjusted network - zero edge nodes deleted

1AYM.1_HBNAO Riconnaviridae 2BUK.A STM Alberovirus FMD Bis 94 Period Control Control 5J98. ACSR THRYMONOUSVIridae 2P17.A Pirin . kaustophilus 2IZW.C_RGNB 6 SSEM SAME DO Circoviridae 1J83.A_CBM1 C.cellulovorans 1JS9.C_BM\ Bromoviridae 1AUYA_TYMO_TygreyRidaepcvo_Cignoxindaelgcvo_dolyomaviridae 4WV8.A_Lectio_V.macrocarpa 3ZMO.A P23.77 Sphaerolipoviridae 1NY7.1 CPMN Secoviridae 1QJZ.B PhMO Tymoviridae 3N35.A Lectin (6.corallodendron 1ZA7.A_CCMO Bromovirida &L.1_BPMO Secoviridae 1NOV.C NOW Nodaviridae 2QE3.A TNB H.sapiens 2BBV.C BBV Nodaviridae 3URF.A_TNO_H,sapiens 1SVA.1 SV40 Polyomaviridae 1STM A SPMOV Papanivirus 1SID.B_MuPVOPolyomaviridae 2TNF.A TNIOM.musculus 2WQ4.A_BclA B.cenocepacia A34.A_STM Virtovirus 2HEY.F_TNB H.sapiens 5IBV.A_HASTIPSISIO TO MUSVIridae 1M06.F_alpha3_Microviridae 2CDO.A CBMO S degradas C.perfringens 2R5I.A HPV18 CONTINUATION OF THE PROPERTY OF T 3M3DC thermocellum U5YA_TNIOM.musculus 1WCD.J.J.B. Birnarinaeparvoviridae 2BPA.1_phiX101_Microviridae 1DNVA_Gmto_Paryexity and Birnaviridae 2W3J.A_CBM55 bacterium CBM/62 CBM32.C Reptitingens 40US. AR 60 Ain BEIRE Boanthracis 3HAG.A_HEO Hepevindae M.musculus 1R64. A Kex Serevisiae illomaviridae 4CRR.A CBM6 Z.galactanivorans.5FUI.A. 1DZL.A HPV16 Papillomaviridae 4D3L.A CBM70 R.flavefaciens 1LP3 A AAVO Parvoviridae 1GMM.A_CBMOR.thermoceNQD.A_CoBD_H.histolytica 3AFG.B_TkSPOT.kodakarensis 3S6P.A 49QSV A Private avindate aviridae 1K5J.B MP X.laevis 2VTX.B_INTYX2abySulto_T.maritima 1MVM.A MVM Parvoviridae 10HF.A Nomega Alphatetraviridae 2OZJ.A_Cup D.hafniense 1XB9.A MP X.laevis

1NLQ.A NP melanogaster

3FJS.A Cupin pinatubonensis