Optimal growth temperature per Archaeal family f_Baldrarchaeaceae (n=1)
f_Borrarchaeaceae (n=5)
f_CR-4 (n=1)
f_CR-4 (n=1)
f_DAOWEDOI (n=1)
f_DXJG01 (n=4)
f_Freyrarchaeaceae (n=29)
f_HEL-GB-A (n=4)
f_HEL-GB-B (n=1)
f_HEL-GB-B (n=1)
f_Hermodarchaeaceae (n=13)
f_Hodarchaeaceae (n=13)
f_JABLTI01 (n=14)
f_JABLTI01 (n=14)
f_JABLTI01 (n=14)
f_JAGGXT01 (n=1)
f_JAJRWK01 (n=3)
f_JOrdarchaeaceae (n=5)
f_Kariarchaeaceae (n=14)
f_LC-3 (n=2)
f_MK-D1 (n=12)
f_MK-D1 (n=12)
f_Odinarchaeaceae (n=3)
f_RDOG01 (n=2)
f_Sifarchaeaceae (n=3)
f_SI46-22 (n=2)
f_Sigynarchaeaceae (n=1)
f_Sigynarchaeaceae (n=60)
f_Wukongarchaeaceae (n=1)
f_YT1-039 (n=1)
f_YT1-057 (n=1) f LFW-46 (n=2) f MCBC01 (n=1) f Micrarchaeaceae (n=98) f Naiadarchaeaceae (n=1) f Nanoanaerosalinaceae (n=6) f Nanoanaerosalinaceae (n=6) f Nanohalakaliarchaeaceae (n=9) f Nanopusillaceae (n=9) f Nanosalinaceae (n=19) f Norongarragalinaceae (n=19) f Morongarragalinaceae (n=10) f Morongarragalinaceae (n=19) f Morongarragalinaceae (n=19) f Morongarragalinaceae (n=19) f Morongarragalinaceae (n=19) f Morongarragalinaceae (n=10) f Morongarragalinaceae (n=10 Temperature >80 60 45 30 15 f_UBA998 (n=5)
f_UBA9989 (n=1)
f_Undinarchaeaceae (n=3)
f_VGJJ01 (n=5)
f_VGKX01 (n=2)
f_VGLX01 (n=3)
f_WALA01 (n=2)
f_WALL01 (n=5)
f_WALR01 (n=1)
f_WAMD01 (n=3)
f_WAMD01 (n=3)
f_WAPJ01 (n=2)
f_WAPJ01 (n=2)
f_WJEV01 (n=2)
f_WJKC01 (n=3)
f_WJKC01 (n=3)
f_WJKC01 (n=3)
f_WJKC01 (n=3)
f_WJKC01 (n=8)
f_WJKC01 (n=8)
f_WJKC01 (n=6)
f_WJKV01 (n=6)
f_WJKV01 (n=6)
f_WJKV01 (n=6)

0

25

no. of genomes (%)

75

100