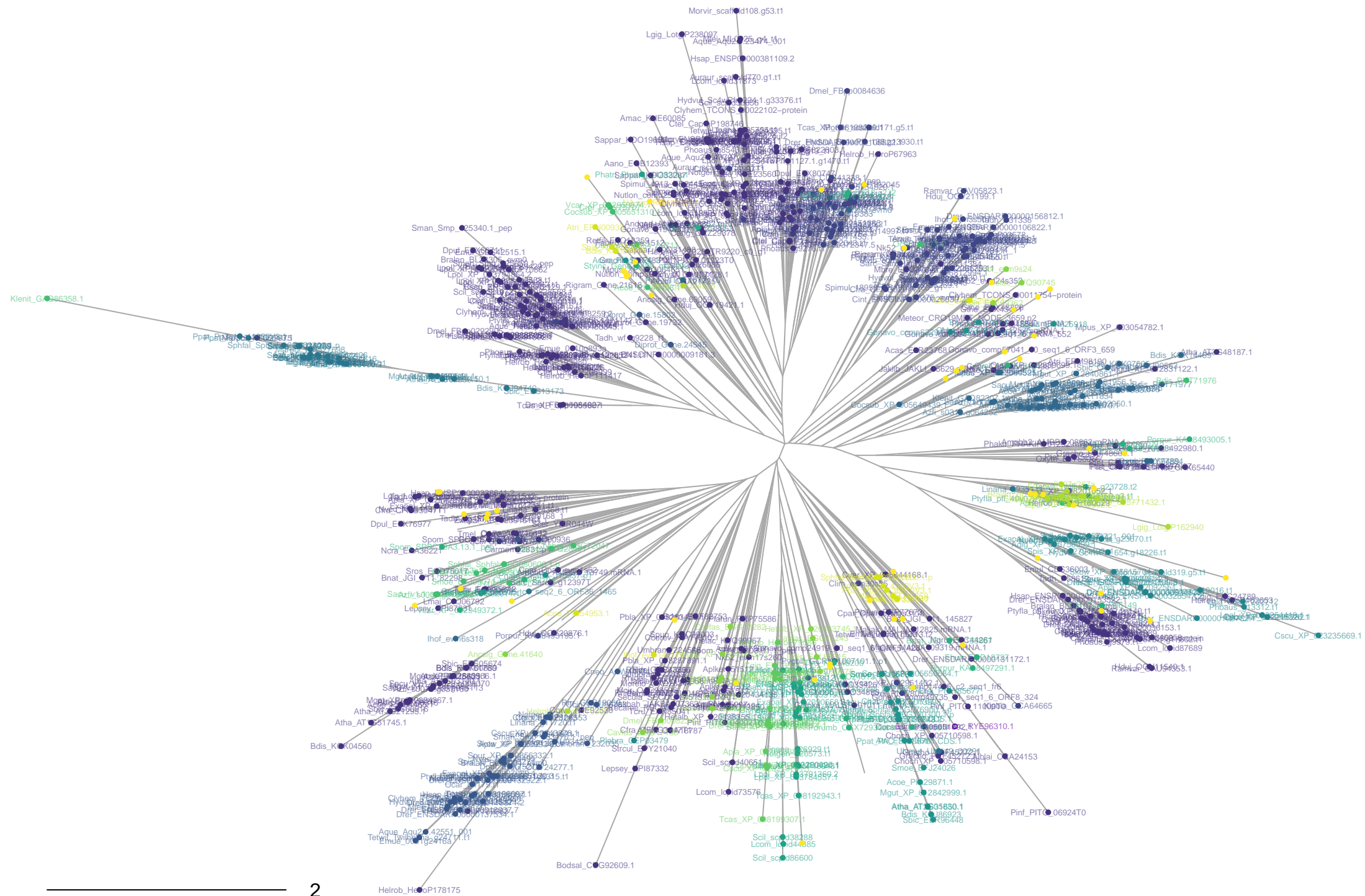


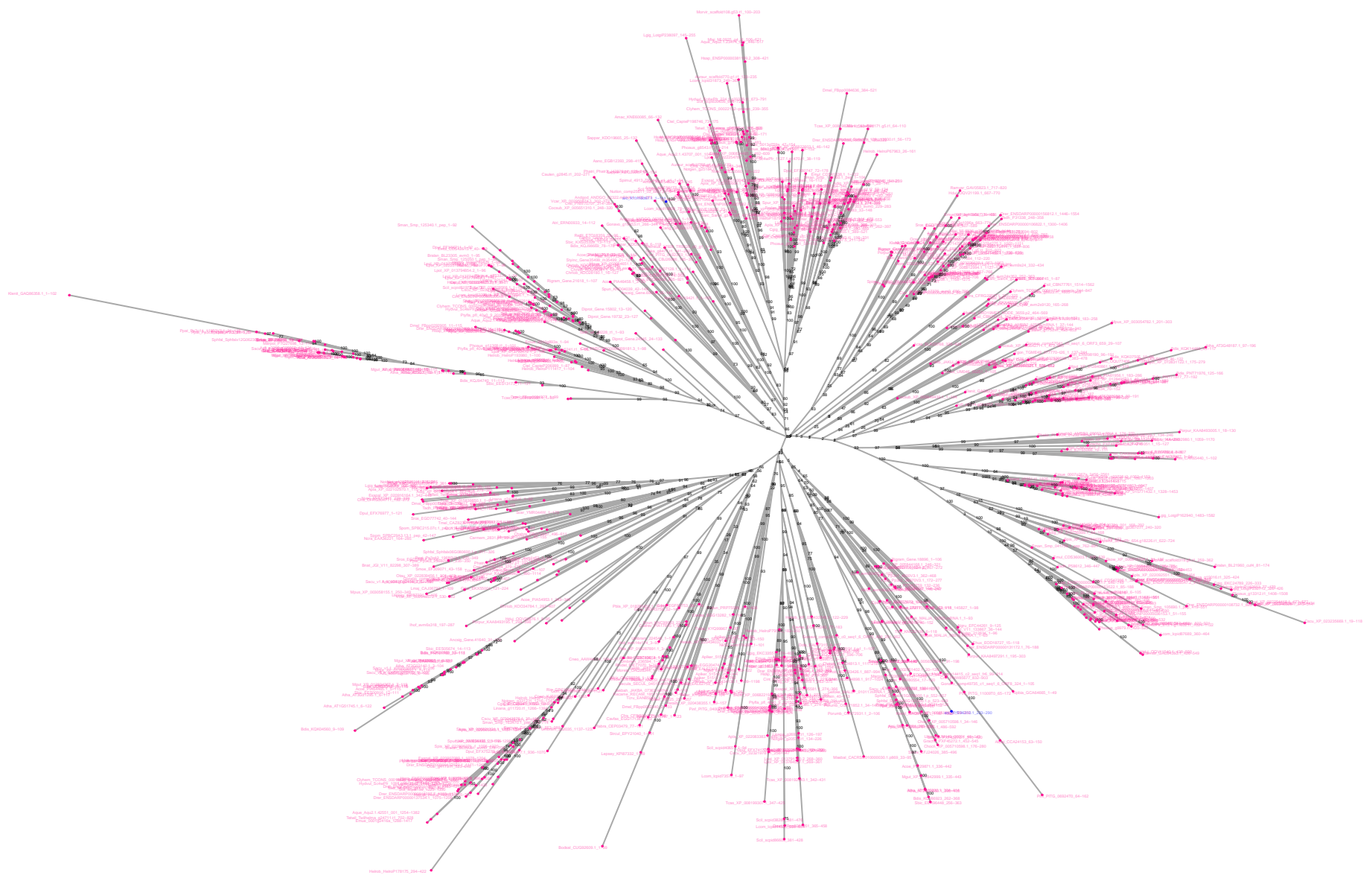
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n=855 sequences

eukaryotes
archaea
bacteria



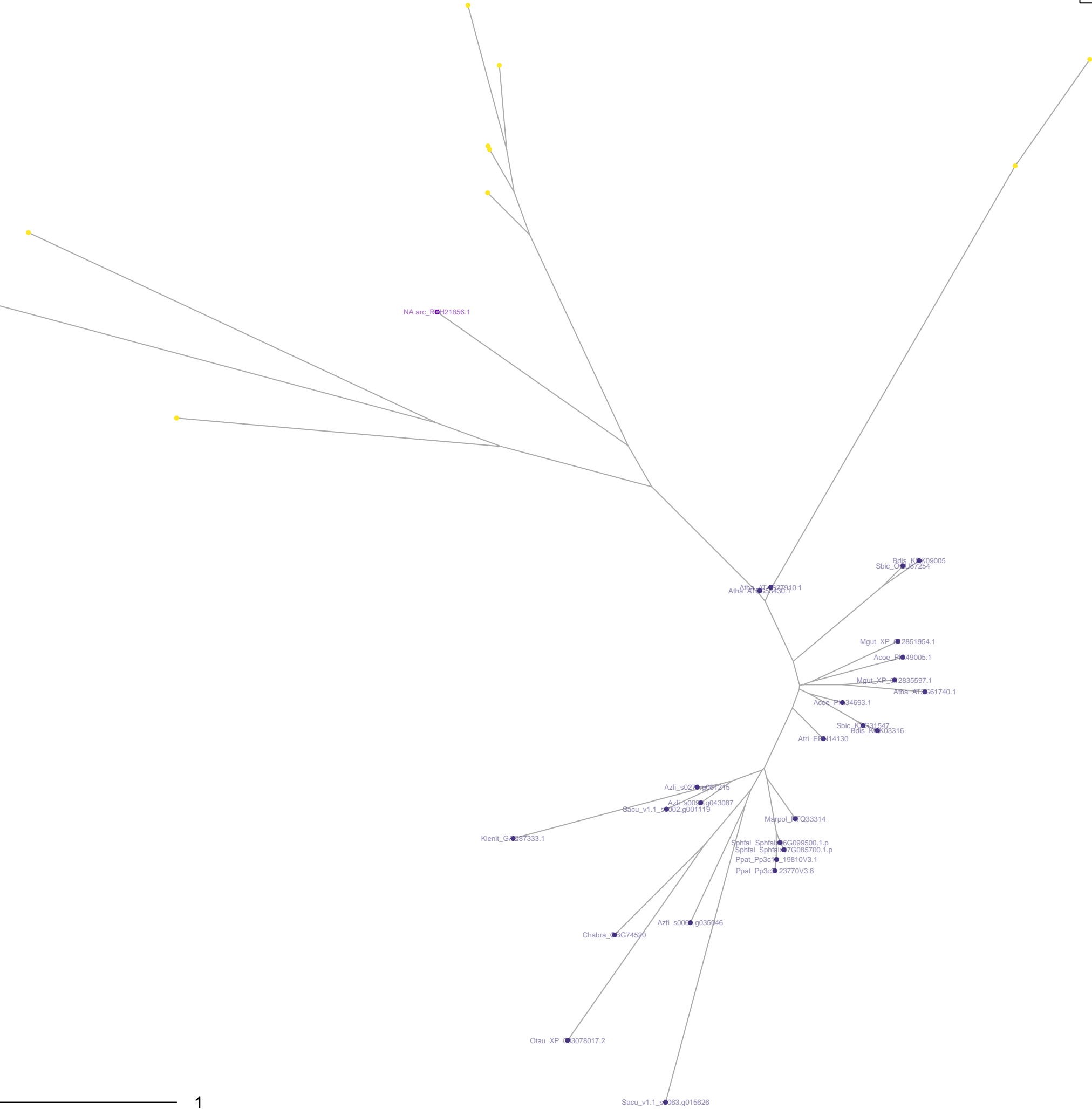
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- PWWP.HG1.2:NSD1/NSD2/NSD3 n = 95
- PWWP.HG1.14:BRD1/BRPF1/BRPF3 n = 49
- PWWP.HG1.20:like:BRD1/BRPF1/BRPF3:likeclu:14 n = 48
- PWWP.HG1.6:like:AC098650.1/BRD1/BRPF1/BRPF3/GLYR1/HDGFL1/HDGFL2/HDGFL3/MSH6/PSIP1/PWWP2A/PWWP2B/PWWP3B/ZCWPW1/ZCWPW2/ZMYND8/ZMYND11:likeclu:8/14/17/18/22 n = 33
- PWWP.HG1.3:DNMT3A/DNMT3B n = 31
- PWWP.HG1.0:like:AC098650.1/BRD1/BRPF1/BRPF3/DNMT3A/DNMT3B/GLYR1/HDGFL1/HDGFL2/HDGFL3/MSH6/NSD1/NSD2/NSD3/PSIP1/PWWP2A/PWWP2B/PWWP3B/ZCWPW1/ZCWPW2/ZMYND8/ZMYND11:likeclu:2/3/8/14/17/18/22 n = 30
- PWWP.HG1.18:ZMYND11 n = 27
- PWWP.HG1.10:like:AC098650.1/BRD1/BRPF1/BRPF3/GLYR1/HDGFL1/HDGFL2/HDGFL3/MSH6/PSIP1/PWWP2A/PWWP2B/PWWP3B/ZCWPW1/ZCWPW2/ZMYND8/ZMYND11:likeclu:8/14/17/18/22 n = 20
- PWWP.HG1.1:like:AC098650.1/BRD1/BRPF1/BRPF3/DNMT3A/DNMT3B/GLYR1/HDGFL1/HDGFL2/HDGFL3/MSH6/NSD1/NSD2/NSD3/PSIP1/PWWP2A/PWWP2B/PWWP3B/ZCWPW1/ZCWPW2/ZMYND8/ZMYND11:likeclu:2/3/8/14/17/18/22 n = 18
- PWWP.HG1.17:ZMYND8 n = 17
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- PWWP.HG1.22:MBD5 n = 14
- PWWP.HG1.12:like:BRD1/BRPF1/BRPF3:likeclu:14 n = 13
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n=36 sequences

eukaryotes
archaea
bacteria



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other n = 9

NA n = 1

