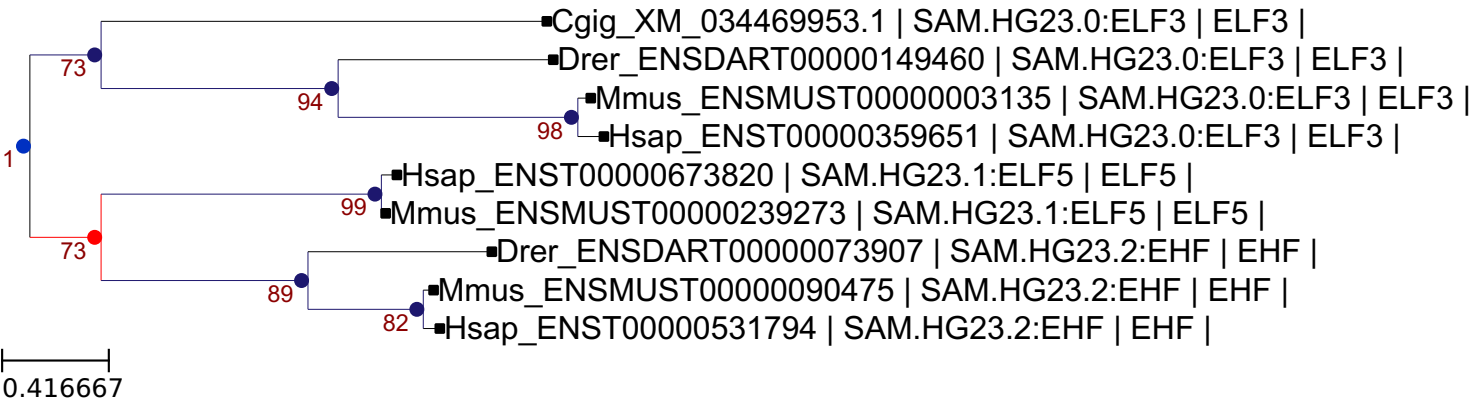
Cgig_XM_034469953.1 | SAM.HG23.0:ELF3 | ELF3 |
Drer_ENSDART00000149460 | SAM.HG23.0:ELF3 | ELF3 |
Mmus_ENSMUST00000003135 | SAM.HG23.0:ELF3 | ELF3 |
Hsap_ENST00000359651 | SAM.HG23.0:ELF3 | ELF3 |
Hsap_ENST00000673820 | SAM.HG23.1:ELF5 | ELF5 |
Mmus_ENSMUST00000239273 | SAM.HG23.1:ELF5 | ELF5 |
Drer_ENSDART00000073907 | SAM.HG23.2:EHF | EHF |
Mmus_ENSMUST00000090475 | SAM.HG23.2:EHF | EHF |
Hsap_ENST00000531794 | SAM.HG23.2:EHF | EHF |
73
94
98
1
99
73
89
82
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