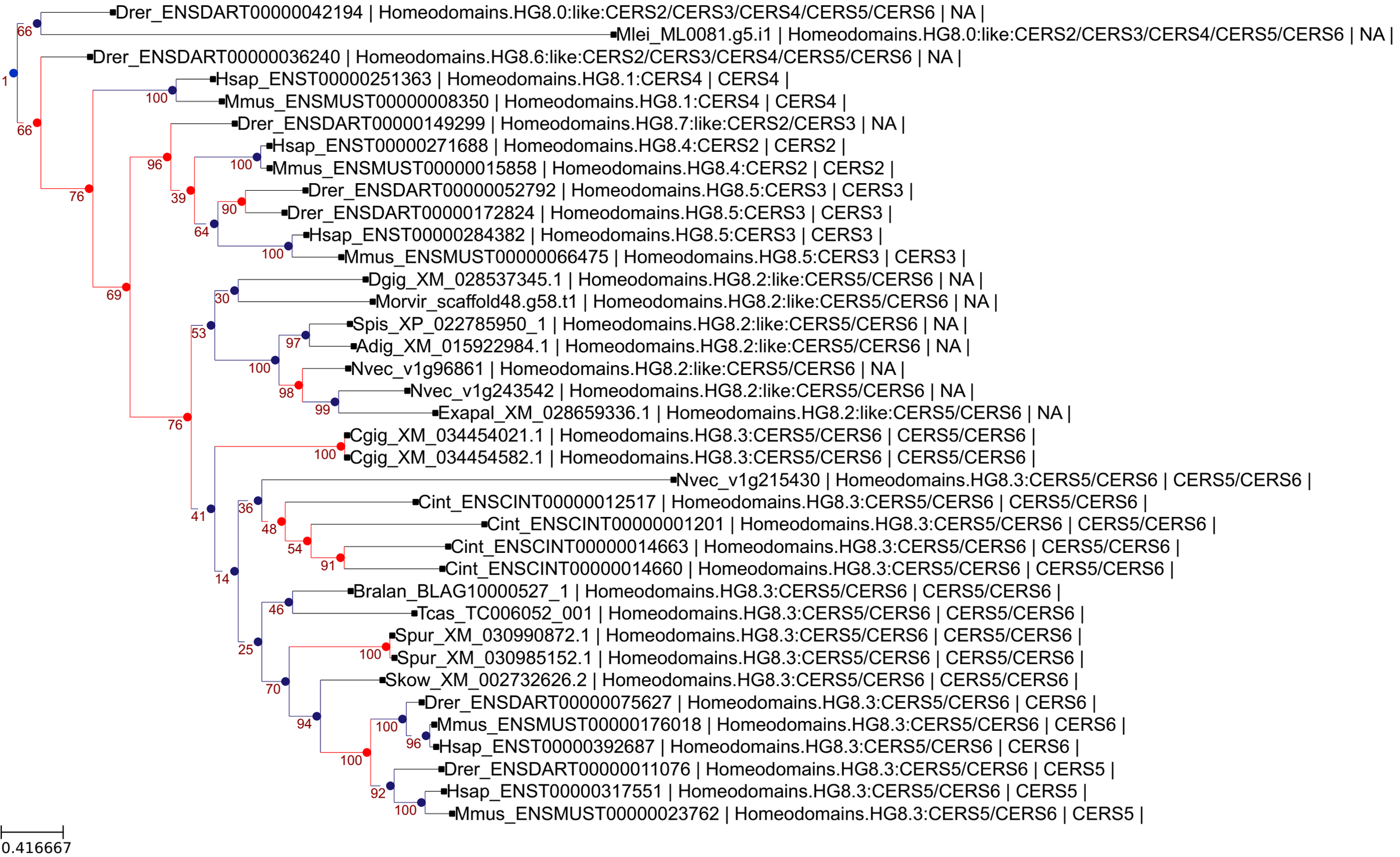
Drer_ENSDART00000042194 | Homeodomains.HG8.0:like:CERS2/CERS3/CERS4/CERS5/CERS6 | NA |
Mlei_ML0081.g5.i1 | Homeodomains.HG8.0:like:CERS2/CERS3/CERS4/CERS5/CERS6 | NA |
Drer_ENSDART00000036240 | Homeodomains.HG8.6:like:CERS2/CERS3/CERS4/CERS5/CERS6 | NA |
Hsap_ENST00000251363 | Homeodomains.HG8.1:CERS4 | CERS4 |
Mmus_ENSMUST00000008350 | Homeodomains.HG8.1:CERS4 | CERS4 |
Drer_ENSDART00000149299 | Homeodomains.HG8.7:like:CERS2/CERS3 | NA |
Hsap_ENST00000271688 | Homeodomains.HG8.4:CERS2 | CERS2 |
Mmus_ENSMUST00000015858 | Homeodomains.HG8.4:CERS2 | CERS2 |
Drer_ENSDART00000052792 | Homeodomains.HG8.5:CERS3 | CERS3 |
Drer_ENSDART00000172824 | Homeodomains.HG8.5:CERS3 | CERS3 |
Hsap_ENST00000284382 | Homeodomains.HG8.5:CERS3 | CERS3 |
Mmus_ENSMUST00000066475 | Homeodomains.HG8.5:CERS3 | CERS3 |
Dgig_XM_028537345.1 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Morvir_scaffold48.g58.t1 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Spis_XP_022785950_1 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Adig_XM_015922984.1 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Nvec_v1g96861 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Nvec_v1g243542 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Exapal_XM_028659336.1 | Homeodomains.HG8.2:like:CERS5/CERS6 | NA |
Cgig_XM_034454021.1 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Cgig_XM_034454582.1 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Nvec_v1g215430 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Cint_ENSCINT00000012517 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Cint_ENSCINT00000001201 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Cint_ENSCINT00000014663 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Cint_ENSCINT00000014660 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Bralan_BLAG10000527_1 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Tcas_TC006052_001 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Spur_XM_030990872.1 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Spur_XM_030985152.1 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Skow_XM_002732626.2 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5/CERS6 |
Drer_ENSDART00000075627 | Homeodomains.HG8.3:CERS5/CERS6 | CERS6 |
Mmus_ENSMUST00000176018 | Homeodomains.HG8.3:CERS5/CERS6 | CERS6 |
Hsap_ENST00000392687 | Homeodomains.HG8.3:CERS5/CERS6 | CERS6 |
Drer_ENSDART00000011076 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5 |
Hsap_ENST00000317551 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5 |
Mmus_ENSMUST00000023762 | Homeodomains.HG8.3:CERS5/CERS6 | CERS5 |
0.416667