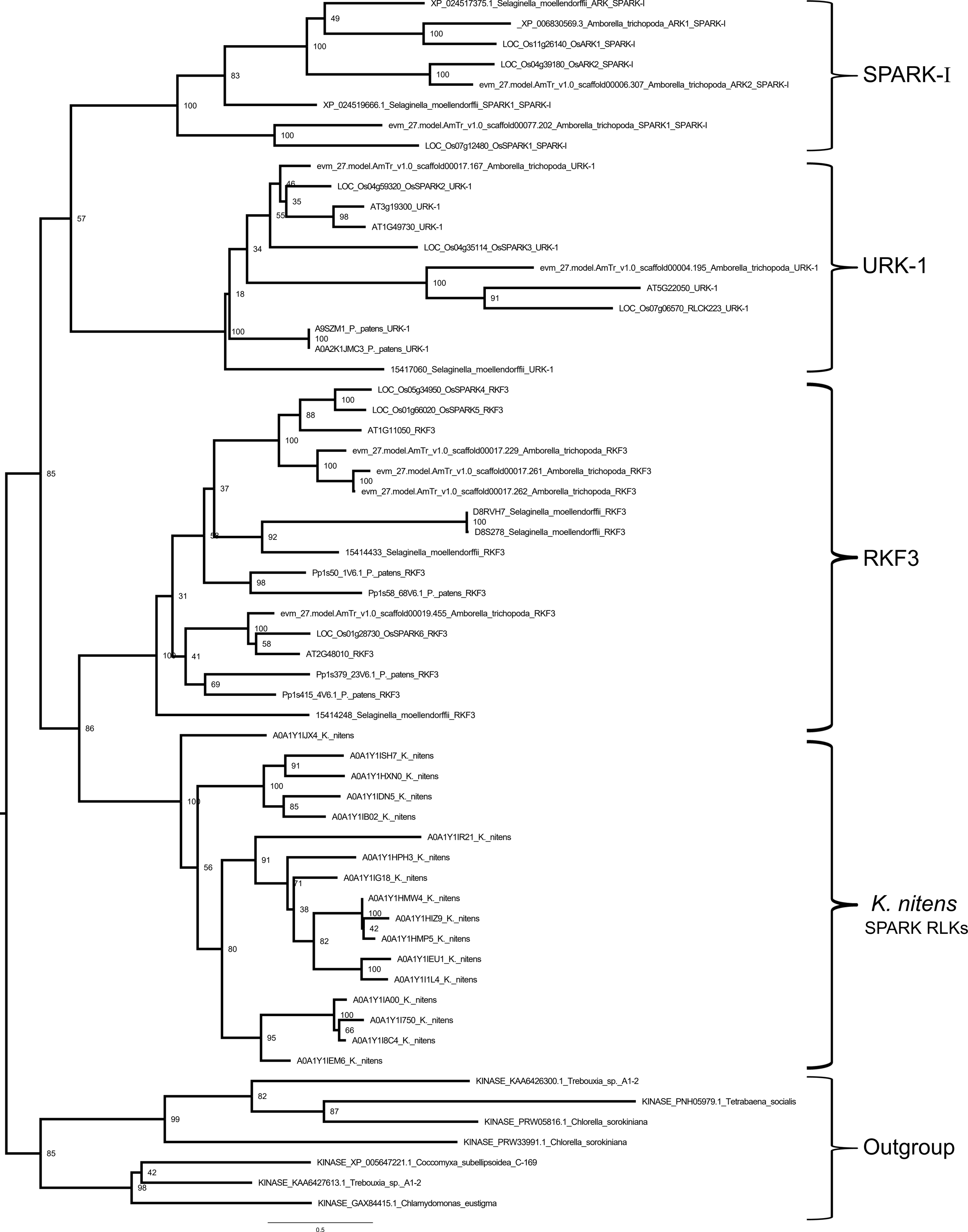
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_XP_006830569.3_Amborella_trichopoda_ARK1_SPARK-I
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LOC_Os04g39180_OsARK2_SPARK-I
evm_27.model.AmTr_v1.0_scaffold00006.307_Amborella_trichopoda_ARK2_SPARK-I
XP_024519666.1_Selaginella_moellendorffii_SPARK1_SPARK-I
evm_27.model.AmTr_v1.0_scaffold00077.202_Amborella_trichopoda_SPARK1_SPARK-I
LOC_Os07g12480_OsSPARK1_SPARK-I
SPARK-I
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LOC_Os04g59320_OsSPARK2_URK-1
AT3g19300_URK-1
AT1G49730_URK-1
LOC_Os04g35114_OsSPARK3_URK-1
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URK-1
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LOC_Os01g66020_OsSPARK5_RKF3
AT1G11050_RKF3
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evm_27.model.AmTr_v1.0_scaffold00017.262_Amborella_trichopoda_RKF3
D8RVH7_Selaginella_moellendorffii_RKF3
D8S278_Selaginella_moellendorffii_RKF3
15414433_Selaginella_moellendorffii_RKF3
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Pp1s58_68V6.1_P._patens_RKF3
evm_27.model.AmTr_v1.0_scaffold00019.455_Amborella_trichopoda_RKF3
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AT2G48010_RKF3
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RKF3
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K. nitens
SPARK RLKs
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KINASE_PNH05979.1_Tetrabaena_socialis
KINASE_PRW05816.1_Chlorella_sorokiniana
KINASE_PRW33991.1_Chlorella_sorokiniana
KINASE_XP_005647221.1_Coccomyxa_subellipsoidea_C-169
KINASE_KAA6427613.1_Trebouxia_sp._A1-2
KINASE_GAX84415.1_Chlamydomonas_eustigma
Outgroup
0.5