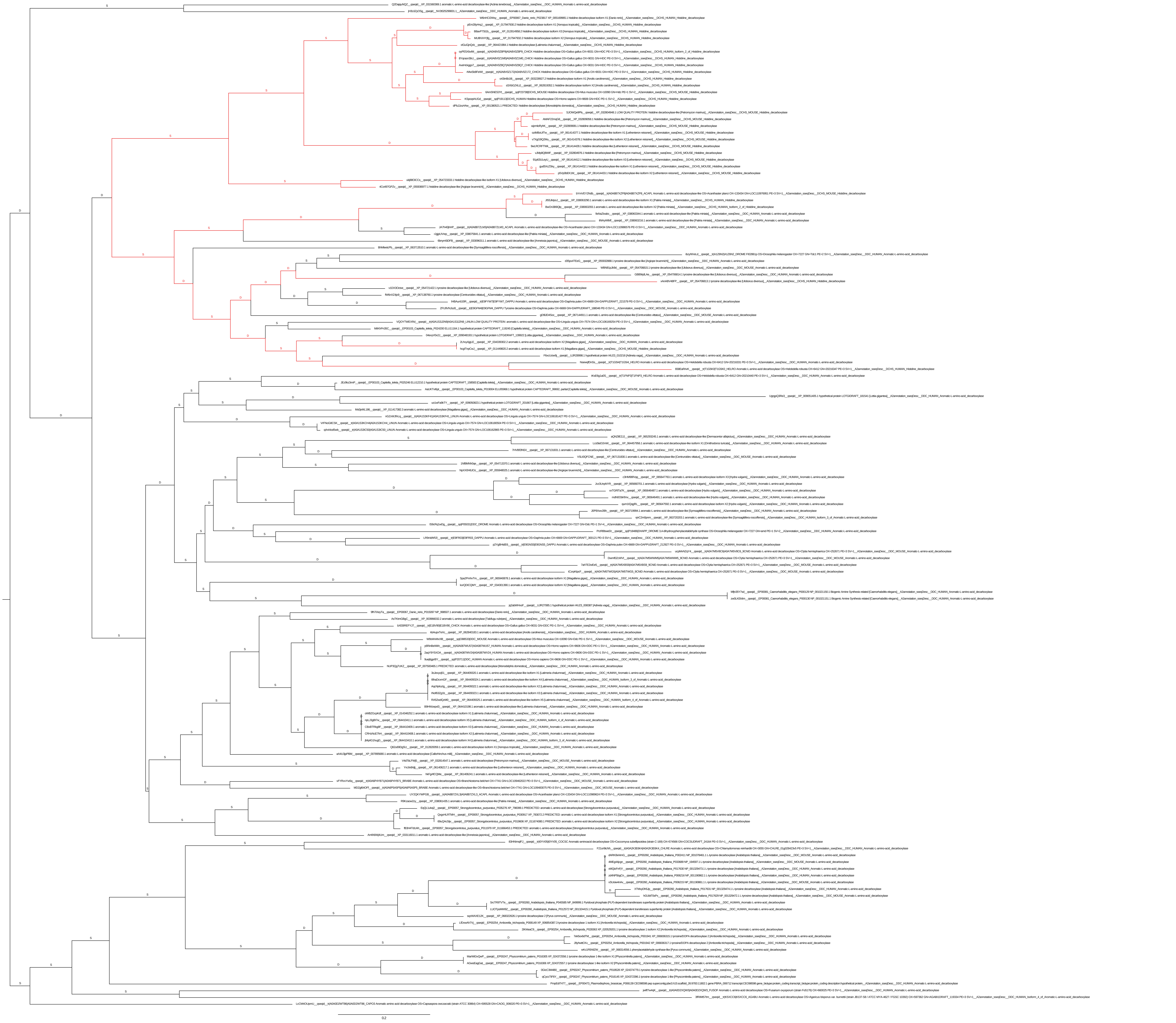
Q2DajquNQZ__qseqid__XP_031560369.1 aromatic-L-amino-acid decarboxylase-like [Actinia tenebrosa]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
jH3L6ZyOSg__qseqid__NV2t025299001.1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
W6nHCl2Wsy__qseqid__EP00067_Danio_rerio_P023817 XP_005169965.1 histidine decarboxylase isoform X1 [Danio rerio]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
pEm28yHrqJ__qseqid__XP_017947930.2 histidine decarboxylase isoform X1 [Xenopus tropicalis]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
B8avP7St1b__qseqid__XP_012814958.2 histidine decarboxylase isoform X3 [Xenopus tropicalis]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
MU8hXHYjfg__qseqid__XP_017947932.2 histidine decarboxylase isoform X2 [Xenopus tropicalis]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
xlGuGjnQck__qseqid__XP_064421984.1 histidine decarboxylase [Latimeria chalumnae]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
oyPE5XbvlM__qseqid__tr|A0A8V0Z8P9|A0A8V0Z8P9_CHICK Histidine decarboxylase OS=Gallus gallus OX=9031 GN=HDC PE=3 SV=1__A2annotation_sseqDesc__DCHS_HUMAN_Isoform_2_of_Histidine_decarboxylase
8Yqrasn3bU__qseqid__tr|A0A8V0Z1M0|A0A8V0Z1M0_CHICK Histidine decarboxylase OS=Gallus gallus OX=9031 GN=HDC PE=3 SV=1__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
AveHckggo7__qseqid__tr|A0A8V0Z8Q7|A0A8V0Z8Q7_CHICK Histidine decarboxylase OS=Gallus gallus OX=9031 GN=HDC PE=3 SV=1__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
rMwSb8FeWI__qseqid__tr|A0A8V0Z172|A0A8V0Z172_CHICK Histidine decarboxylase OS=Gallus gallus OX=9031 GN=HDC PE=3 SV=1__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
o43in6b1l6__qseqid__XP_003228927.2 histidine decarboxylase isoform X1 [Anolis carolinensis]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
sSXbGZ4LtJ__qseqid__XP_062819352.1 histidine decarboxylase isoform X2 [Anolis carolinensis]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
6AmSNES3Yt__qseqid__sp|P23738|DCHS_MOUSE Histidine decarboxylase OS=Mus musculus OX=10090 GN=Hdc PE=1 SV=2__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
KSpoqshUGd__qseqid__sp|P19113|DCHS_HUMAN Histidine decarboxylase OS=Homo sapiens OX=9606 GN=HDC PE=1 SV=2__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
dPbJJzuHAw__qseqid__XP_001380521.1 PREDICTED: histidine decarboxylase [Monodelphis domestica]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
SJOWQe9Pls__qseqid__XP_032804948.1 LOW QUALITY PROTEIN: histidine decarboxylase-like [Petromyzon marinus]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
AMAF22mqG6__qseqid__XP_032809058.1 histidine decarboxylase-like [Petromyzon marinus]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
wjzmloRyhK__qseqid__XP_032809081.1 histidine decarboxylase-like [Petromyzon marinus]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
oziMBoUfTw__qseqid__XP_061414377.1 histidine decarboxylase-like isoform X1 [Lethenteron reissneri]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
v7XgG9Q2Mu__qseqid__XP_061414378.1 histidine decarboxylase-like isoform X2 [Lethenteron reissneri]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
9wLRCRFTWk__qseqid__XP_061414428.1 histidine decarboxylase-like [Lethenteron reissneri]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
vJMp8Q8WIF__qseqid__XP_032804976.1 histidine decarboxylase-like [Petromyzon marinus]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
91p63ULeyU__qseqid__XP_061414412.1 histidine decarboxylase-like isoform X3 [Lethenteron reissneri]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
gud5XcZSky__qseqid__XP_061414432.1 histidine decarboxylase-like isoform X1 [Lethenteron reissneri]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
p5Vp0bEK1M__qseqid__XP_061414433.1 histidine decarboxylase-like isoform X2 [Lethenteron reissneri]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
ukj88OiCCs__qseqid__XP_054723333.1 histidine decarboxylase-like isoform X1 [Uloborus diversus]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
4Co497GPZv__qseqid__XP_055936977.1 histidine decarboxylase-like [Argiope bruennichi]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
bYmVEY2Ndb__qseqid__tr|A0A8B7XZP8|A0A8B7XZP8_ACAPL Aromatic-L-amino-acid decarboxylase-like OS=Acanthaster planci OX=133434 GN=LOC110976951 PE=3 SV=1__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
JI55JkIpoJ__qseqid__XP_038063290.1 aromatic-L-amino-acid decarboxylase-like isoform X1 [Patiria miniata]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
BwOn3B9Qlp__qseqid__XP_038063293.1 aromatic-L-amino-acid decarboxylase-like isoform X2 [Patiria miniata]__A2annotation_sseqDesc__DCHS_HUMAN_Isoform_2_of_Histidine_decarboxylase
9sNaZtxabc__qseqid__XP_038063344.1 aromatic-L-amino-acid decarboxylase-like [Patiria miniata]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
ihlAyWtMf__qseqid__XP_038063216.1 aromatic-L-amino-acid decarboxylase-like [Patiria miniata]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
JA7h40jhHP__qseqid__tr|A0A8B7ZLM3|A0A8B7ZLM3_ACAPL Aromatic-L-amino-acid decarboxylase-like OS=Acanthaster planci OX=133434 GN=LOC110986576 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
rJgjpUVrep__qseqid__XP_038075841.1 aromatic-L-amino-acid decarboxylase-like [Patiria miniata]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
6bnym5iDFB__qseqid__XP_033096311.1 aromatic-L-amino-acid decarboxylase-like [Anneissia japonica]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
BhMleetcPb__qseqid__XP_063713510.1 aromatic-L-amino-acid decarboxylase-like [Symsagittifera roscoffensis]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
ibzyhlVeLd__qseqid__tr|A1Z6N2|A1Z6N2_DROME FI02861p OS=Drosophila melanogaster OX=7227 GN=Tdc1 PE=2 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
d35px4TEeG__qseqid__XP_055932888.1 tyrosine decarboxylase-like [Argiope bruennichi]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
WBN81pJk9d__qseqid__XP_054706815.1 tyrosine decarboxylase-like [Uloborus diversus]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
GBB9rjdLAe__qseqid__XP_054706814.1 tyrosine decarboxylase-like [Uloborus diversus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
xAnABV48FP__qseqid__XP_054706813.1 tyrosine decarboxylase-like [Uloborus diversus]__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
v1DX3Octxe__qseqid__XP_054721422.1 tyrosine decarboxylase-like [Uloborus diversus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
fW6nXZ4jo9__qseqid__XP_067138760.1 tyrosine decarboxylase [Centruroides vittatus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
lHBAun0J0R__qseqid__tr|E9FYW7|E9FYW7_DAPPU Aromatic-L-amino-acid decarboxylase OS=Daphnia pulex OX=6669 GN=DAPPUDRAFT_221579 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
ZFU9VNJsz8__qseqid__tr|E9GFM4|E9GFM4_DAPPU Tyrosine decarboxylase OS=Daphnia pulex OX=6669 GN=DAPPUDRAFT_188046 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
gD9lJD4Soc__qseqid__XP_067144911.1 aromatic-L-amino-acid decarboxylase-like [Centruroides vittatus]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
VQOYTWEXRd__qseqid__tr|A0A1S3J2N9|A0A1S3J2N9_LINUN LOW QUALITY PROTEIN: aromatic-L-amino-acid decarboxylase-like OS=Lingula unguis OX=7574 GN=LOC106169254 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
MiIKhPn35C__qseqid__EP00103_Capitella_teleta_P024200 ELU11164.1 hypothetical protein CAPTEDRAFT_119245 [Capitella teleta]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
04evyV0x2J__qseqid__XP_009048193.1 hypothetical protein LOTGIDRAFT_139922 [Lottia gigantea]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
2LhvyAjgLE__qseqid__XP_034339302.2 aromatic-L-amino-acid decarboxylase isoform X2 [Magallana gigas]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
hcgITnpCwJ__qseqid__XP_011449820.2 aromatic-L-amino-acid decarboxylase isoform X1 [Magallana gigas]__A2annotation_sseqDesc__DCHS_MOUSE_Histidine_decarboxylase
F6xcUciw9j__qseqid__UJR28998.1 hypothetical protein I4U23_010216 [Adineta vaga]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
NsewqfDnSs__qseqid__tr|T1G5I4|T1G5I4_HELRO Aromatic-L-amino-acid decarboxylase OS=Helobdella robusta OX=6412 GN=20216331 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
658EaRrivK__qseqid__tr|T1G5K0|T1G5K0_HELRO Aromatic-L-amino-acid decarboxylase OS=Helobdella robusta OX=6412 GN=20216347 PE=3 SV=1__A2annotation_sseqDesc__DCHS_HUMAN_Histidine_decarboxylase
tKsE6g1a0S__qseqid__tr|T1FNP3|T1FNP3_HELRO Aromatic-L-amino-acid decarboxylase OS=Helobdella robusta OX=6412 GN=20210440 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
JEz9kz3nnP__qseqid__EP00103_Capitella_teleta_P025246 ELU12210.1 hypothetical protein CAPTEDRAFT_158583 [Capitella teleta]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
AaUKTvl6pt__qseqid__EP00103_Capitella_teleta_P019004 ELU05968.1 hypothetical protein CAPTEDRAFT_99692, partial [Capitella teleta]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
UgrgpQ3Re3__qseqid__XP_009051405.1 hypothetical protein LOTGIDRAFT_181541 [Lottia gigantea]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
uo1wFa9kTY__qseqid__XP_009050823.1 hypothetical protein LOTGIDRAFT_201667 [Lottia gigantea]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
Ms0jxML186__qseqid__XP_011417382.3 aromatic-L-amino-acid decarboxylase [Magallana gigas]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
nGZxWJfnLq__qseqid__tr|A0A1S3KF41|A0A1S3KF41_LINUN Aromatic-L-amino-acid decarboxylase OS=Lingula unguis OX=7574 GN=LOC106181427 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
VrFNoG8CSK__qseqid__tr|A0A1S3KCH4|A0A1S3KCH4_LINUN Aromatic-L-amino-acid decarboxylase OS=Lingula unguis OX=7574 GN=LOC106180504 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
qAnI4od5wb__qseqid__tr|A0A1S3IC50|A0A1S3IC50_LINUN Aromatic-L-amino-acid decarboxylase OS=Lingula unguis OX=7574 GN=LOC106162865 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
aQNZilE2JJ__qseqid__XP_065293245.1 aromatic-L-amino-acid decarboxylase-like [Dermacentor albipictus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
Lcs5teD2mW__qseqid__XP_064457956.1 aromatic-L-amino-acid decarboxylase-like isoform X1 [Ornithodoros turicata]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
7HVBf2lN5X__qseqid__XP_067131831.1 aromatic-L-amino-acid decarboxylase-like [Centruroides vittatus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
VSLI0QFCNE__qseqid__XP_067131830.1 aromatic-L-amino-acid decarboxylase-like [Centruroides vittatus]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
1Nf8MNh0qe__qseqid__XP_054713370.1 aromatic-L-amino-acid decarboxylase-like [Uloborus diversus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
NpXX0H6JOz__qseqid__XP_055948025.1 aromatic-L-amino-acid decarboxylase-like [Argiope bruennichi]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
c3HM96fVqq__qseqid__XP_065647763.1 aromatic-L-amino-acid decarboxylase isoform X3 [Hydra vulgaris]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
Jvz3UnpNYR__qseqid__XP_065660701.1 aromatic-L-amino-acid decarboxylase [Hydra vulgaris]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
xxTGRlTa7K__qseqid__XP_065646487.1 aromatic-L-amino-acid decarboxylase [Hydra vulgaris]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
mdN0ObKfmv__qseqid__XP_065646491.1 aromatic-L-amino-acid decarboxylase-like [Hydra vulgaris]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
qum1IQqgRL__qseqid__XP_065647592.1 aromatic-L-amino-acid decarboxylase isoform X2 [Hydra vulgaris]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
J0P8Xwv2Bh__qseqid__XP_063719994.1 aromatic-L-amino-acid decarboxylase-like [Symsagittifera roscoffensis]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
vjnC2m0pnm__qseqid__XP_063720203.1 aromatic-L-amino-acid decarboxylase-like [Symsagittifera roscoffensis]__A2annotation_sseqDesc__DDC_HUMAN_Isoform_3_of_Aromatic-L-amino-acid_decarboxylase
lS9cRq1wDg__qseqid__sp|P05031|DDC_DROME Aromatic-L-amino-acid decarboxylase OS=Drosophila melanogaster OX=7227 GN=Ddc PE=1 SV=4__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
PrzRBbweDr__qseqid__sp|P18486|DHAPP_DROME 3,4-dihydroxyphenylacetaldehyde synthase OS=Drosophila melanogaster OX=7227 GN=amd PE=1 SV=2__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
LR9r4dhR2l__qseqid__tr|E9FR03|E9FR03_DAPPU Aromatic-L-amino-acid decarboxylase OS=Daphnia pulex OX=6669 GN=DAPPUDRAFT_300121 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
p2YgBHtdE6__qseqid__tr|E9GNS5|E9GNS5_DAPPU Aromatic-L-amino-acid decarboxylase OS=Daphnia pulex OX=6669 GN=DAPPUDRAFT_212927 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
ucpMvNZqY4__qseqid__tr|A0A7M5V8C6|A0A7M5V8C6_9CNID Aromatic-L-amino-acid decarboxylase OS=Clytia hemisphaerica OX=252671 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
Dwmf0ZcWVI__qseqid__tr|A0A7M5WWM5|A0A7M5WWM5_9CNID Aromatic-L-amino-acid decarboxylase OS=Clytia hemisphaerica OX=252671 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
7ahTE2wEe5__qseqid__tr|A0A7M5XBS9|A0A7M5XBS9_9CNID Aromatic-L-amino-acid decarboxylase OS=Clytia hemisphaerica OX=252671 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
tCzrpKtpcF__qseqid__tr|A0A7M5TWG5|A0A7M5TWG5_9CNID Aromatic-L-amino-acid decarboxylase OS=Clytia hemisphaerica OX=252671 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
Spa2PmhvTm__qseqid__XP_065940878.1 aromatic-L-amino-acid decarboxylase isoform X1 [Magallana gigas]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
kxrQDICQMY__qseqid__XP_034301390.1 aromatic-L-amino-acid decarboxylase isoform X2 [Magallana gigas]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
Mfjn35Y7sd__qseqid__EP00081_Caenorhabditis_elegans_P000129 NP_001021150.1 Biogenic Amine Synthesis related [Caenorhabditis elegans]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
zw0LKEkltm__qseqid__EP00081_Caenorhabditis_elegans_P000130 NP_001021151.1 Biogenic Amine Synthesis related [Caenorhabditis elegans]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
jq2abWHxsF__qseqid__UJR27085.1 hypothetical protein I4U23_008387 [Adineta vaga]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
9fh7iXeyTa__qseqid__EP00067_Danio_rerio_P015097 NP_998507.1 aromatic-L-amino-acid decarboxylase [Danio rerio]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
AsTKtmGBgC__qseqid__XP_003966032.2 aromatic-L-amino-acid decarboxylase [Takifugu rubripes]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
bXEBREFYJ7__qseqid__tr|E1BV90|E1BV90_CHICK Aromatic-L-amino-acid decarboxylase OS=Gallus gallus OX=9031 GN=DDC PE=1 SV=4__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
kbAupxTeXc__qseqid__XP_062840183.1 aromatic-L-amino-acid decarboxylase [Anolis carolinensis]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
W9sWvMxXl8__qseqid__sp|O88533|DDC_MOUSE Aromatic-L-amino-acid decarboxylase OS=Mus musculus OX=10090 GN=Ddc PE=1 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
y6RinBeWkh__qseqid__tr|A0A087WU57|A0A087WU57_HUMAN Aromatic-L-amino-acid decarboxylase OS=Homo sapiens OX=9606 GN=DDC PE=1 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
2aqY9Y5XOA__qseqid__tr|A0A087WV24|A0A087WV24_HUMAN Aromatic-L-amino-acid decarboxylase OS=Homo sapiens OX=9606 GN=DDC PE=1 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
9uej6gprBY__qseqid__sp|P20711|DDC_HUMAN Aromatic-L-amino-acid decarboxylase OS=Homo sapiens OX=9606 GN=DDC PE=1 SV=2__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
NUP3Qg7UKZ__qseqid__XP_007500465.1 PREDICTED: aromatic-L-amino-acid decarboxylase [Monodelphis domestica]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
3oJexyoj51__qseqid__XP_064409320.1 aromatic-L-amino-acid decarboxylase-like isoform X1 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
I8haDcxmGF__qseqid__XP_064409324.1 aromatic-L-amino-acid decarboxylase-like isoform X4 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Isoform_3_of_Aromatic-L-amino-acid_decarboxylase
AqcNj4ozlg__qseqid__XP_064409322.1 aromatic-L-amino-acid decarboxylase-like isoform X2 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
INof63Zg1b__qseqid__XP_064409323.1 aromatic-L-amino-acid decarboxylase-like isoform X3 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
RA52wdQzM0__qseqid__XP_064409325.1 aromatic-L-amino-acid decarboxylase-like isoform X5 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Isoform_4_of_Aromatic-L-amino-acid_decarboxylase
B9HWzepxl3__qseqid__XP_064410198.1 aromatic-L-amino-acid decarboxylase-like [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
oWBZGcpKdf__qseqid__XP_014346252.1 aromatic-L-amino-acid decarboxylase isoform X1 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
npLJ3gBI7w__qseqid__XP_064410411.1 aromatic-L-amino-acid decarboxylase isoform X5 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Isoform_4_of_Aromatic-L-amino-acid_decarboxylase
CBxBTRkg8F__qseqid__XP_064410409.1 aromatic-L-amino-acid decarboxylase isoform X3 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
CRHzNcE7hH__qseqid__XP_064410408.1 aromatic-L-amino-acid decarboxylase isoform X2 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
jMq4G1hugD__qseqid__XP_064410410.1 aromatic-L-amino-acid decarboxylase isoform X4 [Latimeria chalumnae]__A2annotation_sseqDesc__DDC_HUMAN_Isoform_3_of_Aromatic-L-amino-acid_decarboxylase
Q62u69DgSU__qseqid__XP_012820059.1 aromatic-L-amino-acid decarboxylase isoform X1 [Xenopus tropicalis]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
aA4U3jpPBM__qseqid__XP_007895680.1 aromatic-L-amino-acid decarboxylase [Callorhinchus milii]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
V4d7bLFWj5__qseqid__XP_032814547.1 aromatic-L-amino-acid decarboxylase [Petromyzon marinus]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
YvUlv8Ajlj__qseqid__XP_061406217.1 aromatic-L-amino-acid decarboxylase-like [Lethenteron reissneri]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
hkFgAfCQMe__qseqid__XP_061406241.1 aromatic-L-amino-acid decarboxylase-like [Lethenteron reissneri]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
vFYRvvYw5q__qseqid__tr|A0A6P4YB71|A0A6P4YB71_BRABE Aromatic-L-amino-acid decarboxylase OS=Branchiostoma belcheri OX=7741 GN=LOC109462022 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
MDZgl64OFf__qseqid__tr|A0A6P5A5P5|A0A6P5A5P5_BRABE Aromatic-L-amino-acid decarboxylase-like OS=Branchiostoma belcheri OX=7741 GN=LOC109483075 PE=3 SV=1__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
UY2QKYWPGB__qseqid__tr|A0A8B7ZXL3|A0A8B7ZXL3_ACAPL Aromatic-L-amino-acid decarboxylase OS=Acanthaster planci OX=133434 GN=LOC110989624 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
R9Kzaow2zy__qseqid__XP_038061435.1 aromatic-L-amino-acid decarboxylase-like [Patiria miniata]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
EqQL1ulwj2__qseqid__EP00057_Strongylocentrotus_purpuratus_P035276 XP_798399.1 PREDICTED: aromatic-L-amino-acid decarboxylase [Strongylocentrotus purpuratus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
QxgvHLRTMH__qseqid__EP00057_Strongylocentrotus_purpuratus_P030917 XP_783072.2 PREDICTED: aromatic-L-amino-acid decarboxylase isoform X1 [Strongylocentrotus purpuratus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
69uQAcSijv__qseqid__EP00057_Strongylocentrotus_purpuratus_P019606 XP_011674080.1 PREDICTED: aromatic-L-amino-acid decarboxylase isoform X2 [Strongylocentrotus purpuratus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
fB3H4TdUAh__qseqid__EP00057_Strongylocentrotus_purpuratus_P011979 XP_011666453.1 PREDICTED: aromatic-L-amino-acid decarboxylase [Strongylocentrotus purpuratus]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
AmfA6Wj4Um__qseqid__XP_033116011.1 aromatic-L-amino-acid decarboxylase-like [Anneissia japonica]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
83HNlmqlFJ__qseqid__tr|I0YV05|I0YV05_COCSC Aromatic-aminoacid decarboxylase OS=Coccomyxa subellipsoidea (strain C-169) OX=574566 GN=COCSUDRAFT_24164 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
F21or9tcNh__qseqid__tr|A0A2K3E6K4|A0A2K3E6K4_CHLRE Aromatic-L-amino-acid decarboxylase OS=Chlamydomonas reinhardtii OX=3055 GN=CHLRE_01g028423v5 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
dARK0kAImG__qseqid__EP00260_Arabidopsis_thaliana_P002411 NP_001078461.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
4MEg44jcgn__qseqid__EP00260_Arabidopsis_thaliana_P033689 NP_194597.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
xMQtzFnFjY__qseqid__EP00260_Arabidopsis_thaliana_P017630 NP_001329473.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
zd4NP5bgCn__qseqid__EP00260_Arabidopsis_thaliana_P006216 NP_001190862.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
vSUoiw4nAv__qseqid__EP00260_Arabidopsis_thaliana_P006215 NP_001190861.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
XTMxyDh5Jp__qseqid__EP00260_Arabidopsis_thaliana_P017631 NP_001329474.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
hGUbtT0sPx__qseqid__EP00260_Arabidopsis_thaliana_P017629 NP_001329472.1 L-tyrosine decarboxylase [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
SrcTR9TVTe__qseqid__EP00260_Arabidopsis_thaliana_P045585 NP_849999.1 Pyridoxal phosphate (PLP)-dependent transferases superfamily protein [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
LUOTpsMW9Z__qseqid__EP00260_Arabidopsis_thaliana_P012572 NP_001324415.1 Pyridoxal phosphate (PLP)-dependent transferases superfamily protein [Arabidopsis thaliana]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
sqzWAXE12K__qseqid__XP_068322626.1 tyrosine decarboxylase 2 [Pyrus communis]__A2annotation_sseqDesc__DDC_MOUSE_Aromatic-L-amino-acid_decarboxylase
LfDxwAh7Vj__qseqid__EP00254_Amborella_trichopoda_P008149 XP_006854387.3 tyrosine decarboxylase 1 isoform X1 [Amborella trichopoda]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
2tKIrteaC9__qseqid__EP00254_Amborella_trichopoda_P028363 XP_020529201.1 tyrosine decarboxylase 1 isoform X2 [Amborella trichopoda]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
7ek5ox6dTM__qseqid__EP00254_Amborella_trichopoda_P001941 XP_006836315.1 tyrosine/DOPA decarboxylase 2 [Amborella trichopoda]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
J9yhwttCKv__qseqid__EP00254_Amborella_trichopoda_P001942 XP_006836317.1 tyrosine/DOPA decarboxylase 2 [Amborella trichopoda]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
wKc1R5h9ZW__qseqid__XP_068314556.1 phenylacetaldehyde synthase-like [Pyrus communis]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
MarWtGvQwP__qseqid__EP00247_Physcomitrium_patens_P016305 XP_024372556.1 tyrosine decarboxylase 1-like isoform X1 [Physcomitrella patens]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
AGwsEagGat__qseqid__EP00247_Physcomitrium_patens_P016306 XP_024372557.1 tyrosine decarboxylase 1-like isoform X2 [Physcomitrella patens]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
0GIcC3M4B0__qseqid__EP00247_Physcomitrium_patens_P018528 XP_024374779.1 tyrosine decarboxylase 1-like [Physcomitrella patens]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
qCyccTtF6Y__qseqid__EP00247_Physcomitrium_patens_P016145 XP_024372396.1 tyrosine decarboxylase 1-like [Physcomitrella patens]__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
PmpEdiT477__qseqid__EP00473_Plasmodiophora_brassicae_P006139 CEO98598 pep supercontig:pbe3.h15:scaffold_26:9783:11802:1 gene:PBRA_006712 transcript:CEO98598 gene_biotype:protein_coding transcript_biotype:protein_coding description:hypothetical protein__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
jwiftTw4qK__qseqid__tr|A0A0D2XQW3|A0A0D2XQW3_FUSOF Aromatic-L-amino-acid decarboxylase OS=Fusarium oxysporum (strain Fo5176) OX=660025 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
3fRiMt57tm__qseqid__tr|K5XCC6|K5XCC6_AGABU Aromatic-L-amino-acid decarboxylase OS=Agaricus bisporus var. burnettii (strain JB137-S8 / ATCC MYA-4627 / FGSC 10392) OX=597362 GN=AGABI1DRAFT_119334 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Isoform_4_of_Aromatic-L-amino-acid_decarboxylase
LcCM4OUpmU__qseqid__tr|A0A0D2WT86|A0A0D2WT86_CAPO3 Aromatic amino acid decarboxylase OS=Capsaspora owczarzaki (strain ATCC 30864) OX=595528 GN=CAOG_006020 PE=3 SV=1__A2annotation_sseqDesc__DDC_HUMAN_Aromatic-L-amino-acid_decarboxylase
0.2