

Supplementary Information

for

Unique Loss of the PYHIN Gene Family in Bats Amongst Mammals: Implications for
Inflammasome Sensing

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Supplementary Data S1. PYRIN domain alignment sequences

>Armadillo_AIM2
KEILLKGLDDITDEELQKFKFLLRDVLSSTGKLEKANRTEVADLMIQNAGG
VSALDKTIAIFQKL-NYMHQAKILK-

>Armadillo_MNDA
KKIVLLKGFEAISDYHFNTIKSLLASDLKLTKKMQDDYDRIKIADLMEKKFPG
PTCVDKLIELLKDIPSLKDIVETLRK

>Chimpanzee_AIM2
KEILLTGLDNITDEELDRFKFLLSDEFNIATGKLHTANRIQVANLMIQNAGAV
SAVMKTIRIFQKL-NYMLLAKRLQE

>Chimpanzee_AIM2l
KEILLTSLDNITDEELDRCKCFLPDDFNIATGKLHTGNSTSSQLDLKRWRGV-
-
CSEEDGIFQKL-NYMLVAKCLRE

>Chimpanzee_IFI16
KNIVLLKGLEVINDYHFRMVKSLLSNDLKLNLKMREEYDKIQIADLMEEKFR
GDAGLGKLIQIFKDIPTLEDLAETLKK

>Chimpanzee_MNDA
KKILLKGFELMDDYHFTTIKSLLAYDLGLTTKMQUEEYNRIKITDLMEKKFQG
VACLDKLIELAKDMPSLKNLVNLRK

>Chimpanzee_PYHIN1
KKIVLLKGLEVINDYHFRIVKSLLSNDLKLNPKMKEEYDKIQIADLMEEKFPG
DAGLGKLIIEFFKEIPTLGDLAETLKR

>Cow_1
KKIVLLKGFQHQHINDKCFNMIKSLLAHDCLKLTRKMQDEYDRVKIADLMELKFR
GPDCVDKLIELVKEIDEIKDLAKVLRK

>Dog_1
KKIILLKGLEPINDYQFSIVKSVLARDLGLTANMQEEYNKVKIADLMEEKFPG
VACVNKLIDLFKEMPSLTDIVKQLRN

>Dog_2
KKIILLKGLEPINDYQFSIVKSVLARDLGLTANMQKEYNKVKIADLMEEKFPG
VACVNKLIDLFKEMQDCEDI-----

>Dolphin_1
KRIVLLKGFQHIKDYQFSMIKSLLARDLRLTSKMQDDYDRIKIADLMEVRYPG
AACVEKLIDLIKDIEELKDLAKTLRK

>Elephant_1
KKIVLLKGLEEINDYHFYIVKSLLAHDRLXPKVQEEHDTIKIADLMAEKFPGI
ACVDKLELLKDIKDL-----

>Elephant_2
KKIVLLKGLSINDYQFNTIKSLLARDLQLTKKMQDEYDRIKIADLMEEKFRG
AASVDKLIELFKDMPQLKDLAKSLR-

>Elephant_3
KKILLKGLQPISEYQFNMVKSVLADLQLNKKQAQYGYNKIQIADLMEDKFP
GATCLDILIELFEDIEGLKDLAKTLR-

>Horse_AIM2
KEILLTGLDNITDEELDRFKFLLPDELKVSTAKLENANRTEVANLMIQNVGA
VSAVTKTIHVFQKL-NYMHVAKSLQE

>Horse_2
-

KIVLLKGLEPISAYHFSMVKSLLAHGLQLTAKPQDEYSKIEIAKLMAEKYRGA
TCVDKLIIDLYKDIPELKS LVKNLR-

>Horse_3

-

KIVLLKGLEHINDYYFNVVKSLLAHDLGLTTKMQDEYNRIKIAHLIEEKFQGA
ACVNKLIPELLKDIPE-----

>Horse_4

KKIILLKGLEPINEYQFSMVKFLASDLKLRKAQNEYNRIQIADLMTENFPRA
TCVDKLIELYKDIAELKQLAKTLKN

>Horse_5

KKILLKGLEPISEYHFSIVKSLFAKDLKLTEKAQNEYNRIQIADLMEEFPEG
ACVDALIDLYKDIAELKDLAKTLK-

>Horse_6

KRIVLLKGLEPISEYHFSIVKSLLAHDLKLSRKAQNEYNRIQIADLMEVKFRGA
ACVDALIKLYKDIAELKELAKTLKN

>Human_POP3

KEILLTSLDNITDEELDRFKCFLPDEFNIATGKLHTLNSTS-SQLDLKRWHGV-
-CSEEDRIFQKL-NYMLVAKCLRE

>Human_AIM2

KEILLTGLDNITDEELDRFKFFLSDEFNIATGKLHTANRIQVATLMIQNAGAV
SAVMKTIRIFQKL-NYMMLAKRLQE

>Human_IFI16

KNIVLLKGLEVINDYHFRMVKSLLSNDLKLNLKMREEYDKIQIADLMEEKFR
GDAGLGKLIKIFEDIPTLEDLAETLKK

>Human_MNDA

KKILLKGFELMDDYHFTSIKSLLAYDLGLTTKMQEEYNRIKITDLMEKKFQG
VACLDKLIELAKDMPSLKNLVNNLRK

>Human_PYHIN1

KKIVLLKGLEVINDYHFRIVKSLLSNDLKLNPKMKEEYDKIQIADLMEEKFPG
DAGLGKLIIEFFKEIPTLGDLAETLKR

>Mouse_AI607873

KRIVLLKGLECINKHHFSLFKSLLARDLSLERDNQEKYSTIQIANMMEEKFKP
DAGLGELIEFCEKVPALRKRAEILKK

>Mouse_AIM2

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MLLLTGLDHITEELKRFKYFALTEFQIARSTLDVADRTELADHLIQSAGAAS
AVTKAINIFQKL-NYMHIANALE-

>Mouse_IFI214

KRIVLLTGLMGINDHDFRMVKSLLSKELKLNKMQDEYDRVKIADLMEDKFP
KDAGVVQLIKLYKQIPGLGDIANKLKN

>Mouse_IFI203

KNIVLLKGLNEMEDYQFRTVKSLLRKELKLTCKMQEDYDRIQLADWIEDKFP
KDAGLDKLIKVCEHIKDLKDLAKKLLK-

>Mouse_IFI204

KRIVLLRGLECINKHYFSLFKSLLARDLNLERDNQEYTTIQIANMMEEKFPA
DSGLGKLIAFCEEVPALRKRAEILKK

>Mouse_IFI205

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RLVLLGLECINKHQFNLFKSLMVKDLNLEEDNQEKYTTFQIANMMVKKFPA
DAGLDRLINFCERVPTLKKRAEILKK

>Mouse_MNDA
KRIVLLRGLLECINKHYFSLFKSLLARDLNLERDNQEYTTIQIANMMEEKFPA
DSGLGKLI EFCEEVPALRKRAEILKK
>Mouse_MNDAL
KKIVLLKGL ES MEDYQFRTVKSLLRKELKLTKKLQEDYDRIQLADWMEDKFP
KYAGLDKLIKVCEHIKDLKDLAKKLLK-
>Mouse_PYDC3
KQIVLLSGLEYMNDYNFRALKSLLNHDLKLTKNMQDDYDRIKIADLMEEKFP
EDAGLSKLI EVCEDIPELAARVDILR-
>Mouse_PYDC4
KQIVLLSGLEYMNDYNFRALKSLLNHDLKLTKNMQDDYDRINIADLMEEKFP
EDAGLSKLI EVCEDIPELAARVDILR-
>Mouse_PYDC1
KQIVLLSGLEYMNDYNFRALKSLLNHDLKLTKNMQDDYDRIKIADLMEEKFP
EDAGLSKLI EVCEDIPELADHVDILRK
>Mouse_PYHIN1
KRIVLLTGLMGINDHDFRMVKSLLSKELKLNRMQDQYDRVKIADLMEDKFP
KDAGVDQLIKLYKQIPGLGDIANKLKN
>Opossum_1
---ILLSGLEELQNE DFKKFKFFLSNDCLMMR--
VHNLDRIDLATCMKKQLPCPSFMNKFIEILKKM-NLNETAKKYK-
>Pig_1
KKIILLKGLEKIDEYHFSMIKSLLAHDLELTREMQDNYDRVKIADLMELKFPGI
TCVNILEELLSDMIK-----
>Pig_2
KRIVLLKGFQHMDDYHFKMIKSLLAHDRLRLTRKMQEYDRIKIADLMEEFR
GAACVDKLI ELVKGIDDIKHLVKPLQ-
>Rat_AIM2
--
MLLLTGLDHIT EELKRFKYLALTEFNIPRKT LN IADRTELADQLIQSAGAASA
VAKAISIFQKL-NYMDIAKALE-
>Rat_2
KKIVLMFGLERINDYNFR TLK SLLTHDLKLTQKMQDDYDRIKIADLMEEKFPE
DAGLSKLI EWEGIPEFEDHADTLRK
>Rat_3
KRIVLLKGLEPLSSHHSFLFKSLLASDLKLERHMQEYTKVQIADMMEDEFDP
DAGLGKFIKFCEDLPALRKRAEILKR
>Rat_4
KEIVLIKGLEDMKDYAFRTIKSLLRKELNLTKKMQDDYDRIQLADLLEDKFPQ
DAGLSKLI EVCESIEELKELTDNLKR
>Tasmanian_devil_1

ILLES TELDEEEFKICKYHLKNRGLIIRMEGKGDRTDIAESMKAKFGAVSALK
HITNVFKKMNLNEIVENLQ-
>Parnell's_mustached_bat
QEVLLLTGLDNITDEELDRFKIFFPDEFITPTGKLEYAYRTR--
KPTIKNSSVVS VLMKNI-IFQKL-NYRHVLQKVSE