

NAMPT protein across Species

Description	Species	Per. Identity to human	length	Accession
nicotinamide phosphoribosyltransferase precursor	Homo sapiens	100.00%	491	NP_005737.1
nicotinamide phosphoribosyltransferase isoform X1	Pan troglodytes	99.80%	491	XP_016801130.1
nicotinamide phosphoribosyltransferase	Pongo abelii	98.98%	491	XP_009234172.1
nicotinamide phosphoribosyltransferase	Nomascus leucogenys	98.37%	491	XP_003268195.2
PREDICTED: nicotinamide phosphoribosyltransferase	Macaca fascicularis	97.35%	491	XP_005550528.1
nicotinamide phosphoribosyltransferase	Heterocephalus glaber	97.15%	491	XP_004839733.1
nicotinamide phosphoribosyltransferase	Ptilocolobus tephrosceles	97.15%	491	XP_023038838.1
nicotinamide phosphoribosyltransferase	Rhinolophus ferrumequinum	96.54%	491	XP_032944413.1
nicotinamide phosphoribosyltransferase	Marmota flaviventris	96.95%	491	XP_034491559.1
nicotinamide phosphoribosyltransferase	Sus scrofa	97.15%	491	NP_001026963.1
nicotinamide phosphoribosyltransferase	Fukomys damarensis	96.54%	491	XP_033612645.1
nicotinamide phosphoribosyltransferase	Urocitellus parryii	96.74%	491	XP_026251535.1

PREDICTED: nicotinamide phosphoribosyltransferase	Camelus bactrianus	96.33%	491	XP_010945004.1
nicotinamide phosphoribosyltransferase	Orcinus orca	96.74%	491	XP_004263410.1
nicotinamide phosphoribosyltransferase isoform X1	Octodon degus	96.74%	491	XP_004635047.1
nicotinamide phosphoribosyltransferase	Molossus molossus	96.54%	491	XP_036102529.1
nicotinamide phosphoribosyltransferase	Cebus imitator	96.95%	491	XP_017370928.1
nicotinamide phosphoribosyltransferase isoform X1	Artibeus jamaicensis	96.54%	491	XP_037006380.1
nicotinamide phosphoribosyltransferase isoform X1	Lagenorhynchus obliquidens	96.54%	491	XP_026979692.1
PREDICTED: nicotinamide phosphoribosyltransferase	Hipposideros armiger	96.54%	491	XP_019515705.1
PREDICTED: nicotinamide phosphoribosyltransferase	Chinchilla lanigera	96.33%	491	XP_013365989.1
nicotinamide phosphoribosyltransferase	Sturnira hondurensis	96.54%	491	XP_036912940.1
nicotinamide phosphoribosyltransferase	Callithrix jacchus	96.74%	491	XP_002751770.1
nicotinamide phosphoribosyltransferase	Saimiri boliviensis boliviensis	96.74%	491	XP_010342625.1
nicotinamide phosphoribosyltransferase	Balaenoptera musculus	96.54%	491	XP_036719737.1
nicotinamide phosphoribosyltransferase	Talpa occidentalis	96.74%	491	XP_037367366.1
nicotinamide phosphoribosyltransferase	Desmodus rotundus	96.54%	491	XP_024414335.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Dipodomys ordii	96.33%	491	XP_012890188.1
nicotinamide phosphoribosyltransferase	Phyllostomus discolor	96.33%	491	XP_028380939.1

nicotinamide phosphoribosyltransferase	Tupaia chinensis	96.54%	521	XP_006149233.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X3	Galeopterus variegatus	96.95%	491	XP_008566085.1
PREDICTED: nicotinamide phosphoribosyltransferase	Condylura cristata	96.33%	491	XP_004676980.1
PREDICTED: LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase	Rhinopithecus bieti	96.74%	491	XP_017730166.1
nicotinamide phosphoribosyltransferase	Bos taurus	96.13%	491	NP_001231070.1
PREDICTED: nicotinamide phosphoribosyltransferase	Orycteropus afer afer	96.33%	491	XP_007942627.1
PREDICTED: nicotinamide phosphoribosyltransferase	Sorex araneus	96.33%	491	XP_004602171.1
nicotinamide phosphoribosyltransferase	Pteropus alecto	96.13%	491	XP_006925432.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Aotus nancymaae	96.33%	491	XP_021528844.1
nicotinamide phosphoribosyltransferase	Odocoileus virginianus texanus	96.13%	491	XP_020749167.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Ceratotherium simum simum	96.33%	496	XP_004439352.1
nicotinamide phosphoribosyltransferase isoform X1	Physeter catodon	94.43%	503	XP_028345986.1
nicotinamide phosphoribosyltransferase	Microcebus murinus	96.14%	492	XP_012601885.1
nicotinamide phosphoribosyltransferase-like	Pan paniscus	95.71%	546	XP_034785687.1
nicotinamide phosphoribosyltransferase isoform X2	Ochotona curzoniae	96.13%	495	XP_040857937.1
PREDICTED: nicotinamide phosphoribosyltransferase	Jaculus jaculus	95.93%	491	XP_004652938.1
nicotinamide phosphoribosyltransferase isoform X1	Ochotona curzoniae	96.13%	504	XP_040857936.1

nicotinamide phosphoribosyltransferase isoform X1	Meriones unguiculatus	96.13%	491	XP_021501566.1
nicotinamide phosphoribosyltransferase	Bubalus bubalis	95.93%	491	XP_006042879.2
PREDICTED: nicotinamide phosphoribosyltransferase	Colobus angolensis palliatus	97.04%	553	XP_011812260.1
nicotinamide phosphoribosyltransferase isoform X2	Nannospalax galili	95.93%	491	XP_008828952.1
nicotinamide phosphoribosyltransferase	Rousettus aegyptiacus	96.13%	491	XP_016001357.1
nicotinamide phosphoribosyltransferase	Onychomys torridus	95.93%	491	XP_036061974.1
nicotinamide phosphoribosyltransferase	Suricata suricatta	95.72%	491	XP_029783175.1
nicotinamide phosphoribosyltransferase isoform X1	Pipistrellus kuhlii	95.52%	491	XP_036285604.1
nicotinamide phosphoribosyltransferase isoform X1	Cricetulus griseus	95.72%	491	XP_007647719.1
nicotinamide phosphoribosyltransferase	Mus musculus	95.72%	491	NP_067499.2
nicotinamide phosphoribosyltransferase	Mastomys coucha	95.52%	491	XP_031211570.1
nicotinamide phosphoribosyltransferase isoform X1	Equus caballus	95.72%	491	XP_023494644.1
nicotinamide phosphoribosyltransferase isoform X1	Castor canadensis	95.93%	491	XP_020042025.1
nicotinamide phosphoribosyltransferase	Microtus oregoni	95.72%	491	XP_041514253.1
PREDICTED: nicotinamide phosphoribosyltransferase	Peromyscus maniculatus bairdii	95.52%	491	XP_006989593.1
nicotinamide phosphoribosyltransferase	Hyaena hyaena	95.52%	491	XP_039105714.1
PREDICTED: nicotinamide phosphoribosyltransferase	Oryctolagus cuniculus	95.72%	491	XP_002712079.1

PREDICTED: nicotinamide phosphoribosyltransferase	Erinaceus europaeus	95.93%	491	XP_007527030.1
PREDICTED: nicotinamide phosphoribosyltransferase	Odobenus rosmarus divergens	95.52%	491	XP_004413177.1
nicotinamide phosphoribosyltransferase	Arvicanthis niloticus	95.32%	491	XP_034369958.1
PREDICTED: nicotinamide phosphoribosyltransferase	Marmota marmota marmota	96.83%	501	XP_015343693.1
nicotinamide phosphoribosyltransferase	Grammomys surdaster	95.91%	491	XP_028624622.1
nicotinamide phosphoribosyltransferase isoform X2	Myotis lucifugus	95.32%	491	XP_006085448.1
nicotinamide phosphoribosyltransferase	Ursus arctos horribilis	95.52%	491	XP_026359374.1
nicotinamide phosphoribosyltransferase	Zalophus californianus	95.32%	491	XP_027430885.1
nicotinamide phosphoribosyltransferase	Mesocricetus auratus	95.11%	491	XP_040599209.1
nicotinamide phosphoribosyltransferase	Otolemur garnettii	95.32%	491	XP_012667573.1
PREDICTED: nicotinamide phosphoribosyltransferase	Chrysochloris asiatica	95.11%	491	XP_006834447.1
nicotinamide phosphoribosyltransferase	Rattus norvegicus	95.11%	491	NP_808789.1
PREDICTED: nicotinamide phosphoribosyltransferase	Elephantulus edwardii	96.61%	559	XP_006882748.1
nicotinamide phosphoribosyltransferase	Canis lupus dingo	95.32%	491	XP_025305263.1
nicotinamide phosphoribosyltransferase	Enhydra lutris kenyonii	95.32%	491	XP_022371173.1
nicotinamide phosphoribosyltransferase isoform X1	Sus scrofa	97.25%	478	XP_020957916.1
nicotinamide phosphoribosyltransferase	Neomonachus schauinslandi	95.32%	491	XP_021538607.1

nicotinamide phosphoribosyltransferase	Felis catus	95.11%	491	XP_011278630.1
nicotinamide phosphoribosyltransferase isoform X2	Mus caroli	94.91%	491	XP_021035018.1
nicotinamide phosphoribosyltransferase isoform X1	Myotis myotis	95.11%	491	XP_036183438.1
nicotinamide phosphoribosyltransferase	Mirounga leonina	94.91%	541	XP_034876262.1
nicotinamide phosphoribosyltransferase isoform X1	Trichechus manatus latirostris	95.52%	491	XP_004382662.1
nicotinamide phosphoribosyltransferase	Balaenoptera acutorostrata scammoni	96.61%	520	XP_007180961.1
nicotinamide phosphoribosyltransferase	Leptonychotes weddellii	95.32%	491	XP_030882741.1
nicotinamide phosphoribosyltransferase	Carlito syrichta	96.61%	476	XP_021575127.1
PREDICTED: nicotinamide phosphoribosyltransferase	Propithecus coquereli	95.32%	519	XP_012505782.1
nicotinamide phosphoribosyltransferase	Phoca vitulina	94.70%	491	XP_032258250.1
nicotinamide phosphoribosyltransferase	Arvicola amphibius	94.70%	491	XP_038191738.1
nicotinamide phosphoribosyltransferase isoform X1	Choloepus didactylus	95.32%	491	XP_037692333.1
nicotinamide phosphoribosyltransferase	Sarcophilus harrisii	95.11%	491	XP_031794612.1
nicotinamide phosphoribosyltransferase isoform X1	Myotis lucifugus	93.41%	501	XP_023606846.1
PREDICTED: nicotinamide phosphoribosyltransferase	Miniopterus natalensis	96.40%	522	XP_016060235.1
nicotinamide phosphoribosyltransferase	Phascolarctos cinereus	94.91%	491	XP_020840236.1
nicotinamide phosphoribosyltransferase	Vombatus ursinus	94.91%	491	XP_027694490.1

nicotinamide phosphoribosyltransferase	Trichosurus vulpecula	94.91%	491	XP_036615486.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase	Halichoerus grypus	94.09%	491	XP_035964951.1
nicotinamide phosphoribosyltransferase isoform X1	Eptesicus fuscus	94.30%	491	XP_008148213.1
nicotinamide phosphoribosyltransferase isoform X1	Manis pentadactyla	92.46%	491	XP_036748385.1
nicotinamide phosphoribosyltransferase isoform X1	Manis javanica	92.26%	491	XP_017535371.1
nicotinamide phosphoribosyltransferase isoform X2	Numida meleagris	94.27%	609	XP_021240256.1
nicotinamide phosphoribosyltransferase	Melopsittacus undulatus	94.27%	493	XP_033918755.1
nicotinamide phosphoribosyltransferase	Gallus gallus	94.27%	493	NP_001025899.1
nicotinamide phosphoribosyltransferase isoform X1	Apteryx rowi	94.07%	493	XP_025914795.1
nicotinamide phosphoribosyltransferase isoform X2	Anas platyrhynchos	94.07%	493	XP_027315003.1
nicotinamide phosphoribosyltransferase isoform X2	Falco peregrinus	93.87%	493	XP_005243543.1
nicotinamide phosphoribosyltransferase isoform X2	Aquila chrysaetos chrysaetos	94.07%	493	XP_029870326.1
nicotinamide phosphoribosyltransferase isoform X2	Corvus moneduloides	93.87%	493	XP_031962173.1
nicotinamide phosphoribosyltransferase	Nothoprocta perdicaria	93.87%	493	XP_025902140.1
nicotinamide phosphoribosyltransferase isoform X2	Falco naumanni	93.66%	493	XP_040450378.1
nicotinamide phosphoribosyltransferase isoform X2	Loxodonta africana	93.50%	492	XP_010585648.1
nicotinamide phosphoribosyltransferase	Cavia porcellus	96.19%	483	XP_023420874.1

PREDICTED: nicotinamide phosphoribosyltransferase	Mandrillus leucophaeus	96.95%	571	XP_011841781.1
nicotinamide phosphoribosyltransferase	Ictidomys tridecemlineatus	92.26%	470	XP_040146809.1
nicotinamide phosphoribosyltransferase isoform X2	Lonchura striata domestica	93.66%	493	XP_021390345.1
nicotinamide phosphoribosyltransferase-like	Peromyscus leucopus	96.11%	473	XP_037058088.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X4	Lepidothrix coronata	93.66%	493	XP_017666982.1
nicotinamide phosphoribosyltransferase	Dromaius novaehollandiae	93.66%	493	XP_025953422.1
nicotinamide phosphoribosyltransferase	Echinops telfairi	94.27%	496	XP_004702837.1
nicotinamide phosphoribosyltransferase isoform X1	Numida meleagris	93.13%	615	XP_021240255.1
nicotinamide phosphoribosyltransferase isoform X2	Serinus canaria	93.66%	493	XP_030093868.1
nicotinamide phosphoribosyltransferase	Tyto alba alba	94.03%	583	XP_032852141.1
PREDICTED: nicotinamide phosphoribosyltransferase	Bos mutus	95.79%	478	XP_005901498.1
nicotinamide phosphoribosyltransferase	Microtus ochrogaster	95.77%	564	XP_026634638.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Galeopterus variegatus	94.85%	513	XP_008566081.1
nicotinamide phosphoribosyltransferase isoform X4	Empidonax traillii	93.46%	493	XP_027745883.1
nicotinamide phosphoribosyltransferase isoform X2	Catharus ustulatus	93.46%	493	XP_032914043.1
nicotinamide phosphoribosyltransferase isoform X2	Calypte anna	93.46%	493	XP_030319061.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Sturnus vulgaris	93.46%	493	XP_014724490.1

nicotinamide phosphoribosyltransferase isoform X1	Aquila chrysaetos chrysaetos	92.93%	499	XP_029870325.1
nicotinamide phosphoribosyltransferase	Molothrus ater	93.25%	493	XP_036256842.1
nicotinamide phosphoribosyltransferase isoform X1	Anas platyrhynchos	92.93%	499	XP_027315001.1
nicotinamide phosphoribosyltransferase isoform X1	Strigops habroptila	92.93%	499	XP_030335387.1
nicotinamide phosphoribosyltransferase isoform X2	Coturnix japonica	93.13%	499	XP_015705013.1
nicotinamide phosphoribosyltransferase isoform X1	Gallus gallus	93.13%	499	XP_015131202.1
nicotinamide phosphoribosyltransferase	Tachyglossus aculeatus	92.68%	492	XP_038597142.1
PREDICTED: nicotinamide phosphoribosyltransferase	Bison bison bison	95.03%	585	XP_010839801.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Pseudopodoces humilis	93.25%	493	XP_005526057.1
nicotinamide phosphoribosyltransferase isoform X1	Falco peregrinus	92.73%	499	XP_027651159.1
nicotinamide phosphoribosyltransferase isoform X1	Corvus moneduloides	92.73%	499	XP_031962172.1
nicotinamide phosphoribosyltransferase isoform X1	Falco naumanni	92.53%	499	XP_040450377.1
nicotinamide phosphoribosyltransferase	Ochotona princeps	96.50%	501	XP_036353104.1
nicotinamide phosphoribosyltransferase isoform X1	Nannospalax galili	94.79%	504	XP_029418190.1
PREDICTED: nicotinamide phosphoribosyltransferase	Mustela putorius furo	95.56%	576	XP_012914488.1
nicotinamide phosphoribosyltransferase isoform X2	Parus major	93.05%	493	XP_015483910.1
nicotinamide phosphoribosyltransferase	Puma concolor	95.56%	603	XP_025785586.1

PREDICTED: nicotinamide phosphoribosyltransferase isoform X3	Lepidothrix coronata	92.53%	499	XP_017666981.1
nicotinamide phosphoribosyltransferase	Ornithorhynchus anatinus	92.68%	492	XP_028933518.1
nicotinamide phosphoribosyltransferase isoform X1	Lonchura striata domestica	92.53%	499	XP_021390344.1
nicotinamide phosphoribosyltransferase isoform X2	Hirundo rustica	92.84%	493	XP_039917516.1
nicotinamide phosphoribosyltransferase isoform X1	Serinus canaria	92.53%	499	XP_030093867.1
nicotinamide phosphoribosyltransferase isoform X1	Catharus ustulatus	92.32%	499	XP_032914042.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Sturnus vulgaris	92.32%	499	XP_014724489.1
nicotinamide phosphoribosyltransferase	Vulpes vulpes	94.56%	558	XP_025839856.1
PREDICTED: nicotinamide phosphoribosyltransferase	Equus przewalskii	95.56%	480	XP_008525618.1
nicotinamide phosphoribosyltransferase isoform X3	Empidonax traillii	92.32%	499	XP_027745882.1
nicotinamide phosphoribosyltransferase isoform X1	Monodon monoceros	96.09%	522	XP_029089428.1
PREDICTED: nicotinamide phosphoribosyltransferase	Panthera tigris altaica	95.35%	609	XP_007084032.1
nicotinamide phosphoribosyltransferase isoform X2	Dermochelys coriacea	92.02%	493	XP_038256931.1
nicotinamide phosphoribosyltransferase isoform X1	Calypte anna	92.32%	499	XP_030319054.1
nicotinamide phosphoribosyltransferase isoform X2	Chelonia mydas	92.23%	493	XP_037745077.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Pseudopodoces humilis	92.12%	499	XP_014110660.1
PREDICTED: nicotinamide phosphoribosyltransferase	Myotis brandtii	95.34%	504	XP_014392514.1

nicotinamide phosphoribosyltransferase	Ailuropoda melanoleuca	95.55%	473	XP_011231449.1
nicotinamide phosphoribosyltransferase isoform X2	Chelonoidis abingdonii	91.62%	495	XP_032643208.1
nicotinamide phosphoribosyltransferase isoform X1	Rattus norvegicus	92.43%	500	XP_038967818.1
nicotinamide phosphoribosyltransferase isoform X1	Parus major	91.92%	499	XP_015483909.1
nicotinamide phosphoribosyltransferase isoform X4	Gopherus evgoodei	91.62%	493	XP_030406904.1
nicotinamide phosphoribosyltransferase	Chrysemys picta bellii	92.02%	493	XP_005292422.1
nicotinamide phosphoribosyltransferase	Mauremys reevesii	91.82%	537	XP_039372968.1
nicotinamide phosphoribosyltransferase isoform X1	Mus caroli	94.71%	521	XP_029340374.1
nicotinamide phosphoribosyltransferase isoform X1	Hirundo rustica	91.72%	499	XP_039917515.1
PREDICTED: nicotinamide phosphoribosyltransferase	Crocodylus porosus	92.42%	489	XP_019399104.1
nicotinamide phosphoribosyltransferase	Pogona vitticeps	91.89%	493	XP_020652917.1
PREDICTED: nicotinamide phosphoribosyltransferase	Alligator mississippiensis	92.83%	489	XP_006276691.1
nicotinamide phosphoribosyltransferase	Terrapene carolina triunguis	91.62%	493	XP_024064262.1
PREDICTED: nicotinamide phosphoribosyltransferase	Gavialis gangeticus	92.62%	489	XP_019364965.1
nicotinamide phosphoribosyltransferase	Podarcis muralis	91.82%	493	XP_028603297.1
nicotinamide phosphoribosyltransferase isoform X1	Chelonoidis abingdonii	90.51%	501	XP_032643207.1
nicotinamide phosphoribosyltransferase	Lacerta agilis	91.24%	493	XP_033017623.1

nicotinamide phosphoribosyltransferase isoform X1	Chelonia mydas	91.11%	499	XP_037745076.1
PREDICTED: nicotinamide phosphoribosyltransferase	Latimeria chalumnae	90.36%	490	XP_006006048.1
PREDICTED: LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase	Myotis davidii	94.70%	498	XP_015426696.1
nicotinamide phosphoribosyltransferase	Thamnophis elegans	91.77%	493	XP_032076888.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Zalophus californianus	92.46%	489	XP_035580517.1
nicotinamide phosphoribosyltransferase isoform X3	Gopherus evgoodei	90.51%	499	XP_030406828.1
PREDICTED: nicotinamide phosphoribosyltransferase	Anolis carolinensis	91.77%	493	XP_003221586.1
PREDICTED: nicotinamide phosphoribosyltransferase	Monodelphis domestica	94.71%	489	XP_007504168.1
nicotinamide phosphoribosyltransferase isoform X1	Coturnix japonica	92.78%	506	XP_015704941.1
PREDICTED: nicotinamide phosphoribosyltransferase	Charadrius vociferus	93.12%	566	XP_009881850.1
nicotinamide phosphoribosyltransferase isoform X1	Trachemys scripta elegans	89.11%	509	XP_034622440.1
PREDICTED: nicotinamide phosphoribosyltransferase	Cariama cristata	94.09%	511	XP_009696119.1
nicotinamide phosphoribosyltransferase	Zootoca vivipara	90.80%	493	XP_034983850.1
PREDICTED: nicotinamide phosphoribosyltransferase	Gavia stellata	93.88%	523	XP_009815497.1
nicotinamide phosphoribosyltransferase isoform X2	Phasianus colchicus	93.91%	493	XP_031453889.1
PREDICTED: nicotinamide phosphoribosyltransferase	Apteryx mantelli mantelli	94.08%	494	XP_013799066.1
nicotinamide phosphoribosyltransferase	Egretta garzetta	93.88%	503	XP_009634368.1

PREDICTED: nicotinamide phosphoribosyltransferase	Haliaeetus albicilla	93.88%	484	XP_009914254.1
nicotinamide phosphoribosyltransferase	Crotalus tigris	90.80%	493	XP_039210400.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Anser cygnoides domesticus	93.89%	509	XP_013046206.1
nicotinamide phosphoribosyltransferase	Columba livia	94.09%	520	XP_021151099.1
PREDICTED: nicotinamide phosphoribosyltransferase	Pterocles gutturalis	94.29%	473	XP_010080393.1
PREDICTED: nicotinamide phosphoribosyltransferase	Nipponia nippon	94.08%	505	XP_009461245.1
PREDICTED: nicotinamide phosphoribosyltransferase	Sinocyclocheilus rhinoceros	88.87%	493	XP_016364083.1
PREDICTED: nicotinamide phosphoribosyltransferase	Sinocyclocheilus grahami	88.87%	493	XP_016125789.1
PREDICTED: nicotinamide phosphoribosyltransferase	Sinocyclocheilus anshuiensis	88.87%	493	XP_016303166.1
PREDICTED: nicotinamide phosphoribosyltransferase	Pygoscelis adeliae	93.68%	538	XP_009326645.1
PREDICTED: nicotinamide phosphoribosyltransferase	Calidris pugnax	93.49%	515	XP_014818853.1
nicotinamide phosphoribosyltransferase isoform X2	Apteryx rowi	94.08%	474	XP_025914796.1
PREDICTED: nicotinamide phosphoribosyltransferase	Cuculus canorus	93.67%	486	XP_009555599.1
PREDICTED: nicotinamide phosphoribosyltransferase	Opisthocomus hoazin	94.08%	528	XP_009932082.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Anser cygnoides domesticus	94.08%	474	XP_013046208.1
PREDICTED: nicotinamide phosphoribosyltransferase	Tauraco erythrolophus	94.08%	473	XP_009985471.1
PREDICTED: nicotinamide phosphoribosyltransferase	Aptenodytes forsteri	93.47%	505	XP_019330121.1

nicotinamide phosphoribosyltransferase	Polypterus senegalus	88.59%	491	XP_039617123.1
PREDICTED: nicotinamide phosphoribosyltransferase	Leptosomus discolor	93.87%	507	XP_009959226.1
PREDICTED: nicotinamide phosphoribosyltransferase	Pelecanus crispus	93.87%	473	XP_009478542.1
nicotinamide phosphoribosyltransferase-like	Carassius auratus	88.45%	556	XP_026096182.1
PREDICTED: nicotinamide phosphoribosyltransferase	Dryobates pubescens	93.67%	476	XP_009900570.1
PREDICTED: nicotinamide phosphoribosyltransferase	Phalacrocorax carbo	93.87%	515	XP_009502423.1
nicotinamide phosphoribosyltransferase isoform X1	Manacus vitellinus	93.66%	491	XP_017923533.1
PREDICTED: nicotinamide phosphoribosyltransferase	Apaloderma vittatum	93.87%	473	XP_009874603.1
PREDICTED: nicotinamide phosphoribosyltransferase	Colius striatus	93.66%	473	XP_010205642.1
PREDICTED: nicotinamide phosphoribosyltransferase	Nestor notabilis	93.87%	473	XP_010022729.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Cyprinus carpio	88.66%	493	XP_018936795.1
PREDICTED: nicotinamide phosphoribosyltransferase	Acanthisitta chloris	93.87%	473	XP_009068452.1
PREDICTED: nicotinamide phosphoribosyltransferase	Chaetura pelagica	93.45%	550	XP_010002152.1
nicotinamide phosphoribosyltransferase	Xiphophorus couchianus	87.40%	491	XP_027900246.1
PREDICTED: nicotinamide phosphoribosyltransferase	Buceros rhinoceros silvestris	93.25%	518	XP_010141490.1
nicotinamide phosphoribosyltransferase	Geospiza fortis	93.08%	507	XP_030914892.1
nicotinamide phosphoribosyltransferase isoform X2	Empidonax traillii	93.45%	519	XP_027745881.1

PREDICTED: nicotinamide phosphoribosyltransferase	Chlamydotis macqueenii	93.66%	473	XP_010119715.1
PREDICTED: nicotinamide phosphoribosyltransferase	Merops nubicus	93.45%	473	XP_008940977.1
nicotinamide phosphoribosyltransferase	Scleropages formosus	88.66%	491	XP_018581859.1
nicotinamide phosphoribosyltransferase	Antrostomus carolinensis	93.84%	473	XP_010170836.1
PREDICTED: nicotinamide phosphoribosyltransferase	Fulmarus glacialis	93.66%	473	XP_009583621.1
PREDICTED: nicotinamide phosphoribosyltransferase	Corvus brachyrhynchos	92.32%	495	XP_017601378.1
nicotinamide phosphoribosyltransferase	Carassius auratus	88.45%	493	XP_026091213.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Galeopterus variegatus	95.10%	500	XP_008566083.1
PREDICTED: nicotinamide phosphoribosyltransferase	Phaethon lepturus	93.45%	473	XP_010289066.1
nicotinamide phosphoribosyltransferase	Pimephales promelas	89.08%	546	XP_039516638.1
nicotinamide phosphoribosyltransferase	Athene cunicularia	91.96%	491	XP_026723806.1
PREDICTED: nicotinamide phosphoribosyltransferase	Gekko japonicus	90.43%	492	XP_015265036.1
nicotinamide phosphoribosyltransferase	Pseudonaja textilis	90.43%	493	XP_026557207.1
nicotinamide phosphoribosyltransferase isoform X2	Pipra filicauda	93.66%	526	XP_027607612.1
nicotinamide phosphoribosyltransferase	Xiphophorus maculatus	86.99%	491	XP_005799137.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Lepidothrix coronata	93.66%	589	XP_017666980.1
nicotinamide phosphoribosyltransferase	Pungitius pungitius	88.47%	486	XP_037316532.1

PREDICTED: nicotinamide phosphoribosyltransferase	Cyprinodon variegatus	87.47%	491	XP_015243752.1
nicotinamide phosphoribosyltransferase	Cyprinodon tularosa	87.27%	491	XP_038130647.1
nicotinamide phosphoribosyltransferase	Falco cherrug	92.69%	485	XP_027653450.1
nicotinamide phosphoribosyltransferase	Anguilla anguilla	88.26%	491	XP_035281178.1
nicotinamide phosphoribosyltransferase	Danio rerio	88.24%	493	XP_002661386.1
nicotinamide phosphoribosyltransferase	Xiphophorus hellerii	86.79%	491	XP_032446026.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase	Gasterosteus aculeatus aculeatus	88.47%	486	XP_040029600.1
PREDICTED: nicotinamide phosphoribosyltransferase	Struthio camelus australis	92.90%	527	XP_009664298.1
PREDICTED: nicotinamide phosphoribosyltransferase	Poecilia mexicana	86.79%	491	XP_014826352.1
nicotinamide phosphoribosyltransferase	Dasypus novemcinctus	95.24%	467	XP_004467608.1
nicotinamide phosphoribosyltransferase isoform X2	Manacus vitellinus	92.50%	483	XP_017923535.1
nicotinamide phosphoribosyltransferase isoform X2	Corapipo altera	92.48%	497	XP_027495440.1
nicotinamide phosphoribosyltransferase	Megalops cyprinoides	87.99%	491	XP_036373721.1
PREDICTED: nicotinamide phosphoribosyltransferase	Tinamus guttatus	92.50%	602	XP_010218515.1
nicotinamide phosphoribosyltransferase isoform X1	Chanos chanos	88.02%	491	XP_030638761.1
PREDICTED: nicotinamide phosphoribosyltransferase	Poecilia formosa	86.59%	491	XP_007579252.1
nicotinamide phosphoribosyltransferase isoform X1	Dermochelys coriacea	90.12%	513	XP_038256922.1

nicotinamide phosphoribosyltransferase	Cyclopterus lumpus	86.63%	490	XP_034382304.1
nicotinamide phosphoribosyltransferase	Cyanistes caeruleus	92.08%	521	XP_023774788.1
nicotinamide phosphoribosyltransferase S homeolog isoform X1	Xenopus laevis	87.42%	491	XP_041443472.1
nicotinamide phosphoribosyltransferase	Etheostoma spectabile	88.47%	491	XP_032360541.1
PREDICTED: nicotinamide phosphoribosyltransferase	Poecilia reticulata	86.59%	491	XP_008399189.1
nicotinamide phosphoribosyltransferase isoform X1	Empidonax traillii	92.28%	525	XP_027745880.1
nicotinamide phosphoribosyltransferase S homeolog	Xenopus laevis	87.42%	491	NP_001082484.1
nicotinamide phosphoribosyltransferase	Zonotrichia albicollis	92.28%	518	XP_005482892.1
nicotinamide phosphoribosyltransferase	Oryzias melastigma	86.86%	491	XP_024143704.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus anshuiensis	87.27%	493	XP_016299755.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X1	Lepidothrix coronata	92.48%	595	XP_017666979.1
nicotinamide phosphoribosyltransferase isoform X1	Pipra filicauda	92.48%	595	XP_027607611.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus rhinoceros	87.82%	493	XP_016381148.1
nicotinamide phosphoribosyltransferase	Oryzias latipes	86.65%	491	XP_023808115.1
nicotinamide phosphoribosyltransferase	Esox lucius	87.27%	491	XP_010894778.1
nicotinamide phosphoribosyltransferase isoform X2	Gopherus evgoodei	89.71%	571	XP_030406737.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase	Trematomus bernacchii	86.83%	490	XP_033987178.1

nicotinamide phosphoribosyltransferase	Toxotes jaculatrix	87.06%	491	XP_040885971.1
PREDICTED: nicotinamide phosphoribosyltransferase	Ficedula albicollis	92.45%	601	XP_016158566.1
nicotinamide phosphoribosyltransferase	Pseudochaenichthys georgianus	86.83%	490	XP_033968405.1
nicotinamide phosphoribosyltransferase isoform X2	Chanos chanos	88.02%	490	XP_030638769.1
PREDICTED: nicotinamide phosphoribosyltransferase	Salmo salar	88.47%	491	XP_014008446.1
nicotinamide phosphoribosyltransferase	Acanthochromis polyacanthus	88.61%	491	XP_022058581.1
nicotinamide phosphoribosyltransferase	Fundulus heteroclitus	87.55%	491	XP_012725964.2
nicotinamide phosphoribosyltransferase	Kryptolebias marmoratus	87.06%	491	XP_017270399.1
nicotinamide phosphoribosyltransferase	Archocentrus centrarchus	88.21%	491	XP_030576003.1
nicotinamide phosphoribosyltransferase	Paramormyrops kingsleyae	87.82%	487	XP_023694072.1
PREDICTED: nicotinamide phosphoribosyltransferase	Stegastes partitus	88.40%	491	XP_008277114.1
nicotinamide phosphoribosyltransferase	Sparus aurata	87.42%	486	XP_030295647.1
nicotinamide phosphoribosyltransferase	Cottoperca gobio	86.63%	490	XP_029317606.1
nicotinamide phosphoribosyltransferase	Etheostoma cragini	88.05%	491	XP_034718842.1
PREDICTED: nicotinamide phosphoribosyltransferase	Nothobranchius furzeri	88.42%	545	XP_015813506.1
nicotinamide phosphoribosyltransferase	Rhinatrema bivittatum	89.94%	491	XP_029471331.1
nicotinamide phosphoribosyltransferase	Sander lucioperca	88.05%	491	XP_031140671.1

LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Nannospalax galili	90.02%	486	XP_029424353.1
nicotinamide phosphoribosyltransferase	Alligator sinensis	93.02%	504	XP_025063163.1
nicotinamide phosphoribosyltransferase	Sphaeramia orbicularis	88.84%	491	XP_030004569.1
nicotinamide phosphoribosyltransferase	Seriola dumerili	88.05%	491	XP_022604965.1
nicotinamide phosphoribosyltransferase	Salmo trutta	88.26%	491	XP_029597694.1
PREDICTED: nicotinamide phosphoribosyltransferase	Lepisosteus oculatus	88.61%	491	XP_006633446.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Salmo salar	88.26%	491	XP_014063204.1
nicotinamide phosphoribosyltransferase-like	Carassius auratus	87.06%	493	XP_026065047.1
nicotinamide phosphoribosyltransferase	Microcaecilia unicolor	89.37%	491	XP_030071801.1
nicotinamide phosphoribosyltransferase	Syngnathus acus	86.79%	487	XP_037098455.1
nicotinamide phosphoribosyltransferase	Anabas testudineus	86.45%	491	XP_026218388.1
nicotinamide phosphoribosyltransferase-like	Salvelinus namaycush	88.05%	491	XP_038852314.1
nicotinamide phosphoribosyltransferase	Perca flavescens	87.84%	491	XP_028426151.1
nicotinamide phosphoribosyltransferase	Python bivittatus	92.39%	473	XP_007440298.1
nicotinamide phosphoribosyltransferase	Epinephelus lanceolatus	88.61%	490	XP_033470580.1
nicotinamide phosphoribosyltransferase	Denticeps clupeoides	86.24%	491	XP_028810490.1
nicotinamide phosphoribosyltransferase-like	Oncorhynchus nerka	88.26%	491	XP_029486081.1

nicotinamide phosphoribosyltransferase	Acanthopagrus latus	87.42%	486	XP_036977970.1
nicotinamide phosphoribosyltransferase isoform X3	Chelonia mydas	89.23%	523	XP_037745078.1
nicotinamide phosphoribosyltransferase	Acipenser ruthenus	88.61%	491	XP_033889216.1
nicotinamide phosphoribosyltransferase-like	Carassius auratus	86.85%	493	XP_026121436.1
nicotinamide phosphoribosyltransferase isoform X4	Chelonia mydas	89.23%	519	XP_037745079.1
nicotinamide phosphoribosyltransferase-like	Megalops cyprinoides	87.71%	491	XP_036376169.1
nicotinamide phosphoribosyltransferase	Electrophorus electricus	87.42%	491	XP_026863327.2
nicotinamide phosphoribosyltransferase	Sebastes umbrosus	86.24%	491	XP_037616685.1
nicotinamide phosphoribosyltransferase	Acipenser ruthenus	88.61%	491	XP_033882852.1
PREDICTED: nicotinamide phosphoribosyltransferase	Hippocampus comes	87.00%	491	XP_019721587.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase	Salaria fasciatus	87.04%	490	XP_029968867.1
nicotinamide phosphoribosyltransferase	Notechis scutatus	90.31%	504	XP_026533026.1
nicotinamide phosphoribosyltransferase	Oncorhynchus tshawytscha	88.05%	491	XP_024228758.1
nicotinamide phosphoribosyltransferase	Nematolebias whitei	87.84%	491	XP_037551853.1
nicotinamide phosphoribosyltransferase	Cheilinus undulatus	87.84%	491	XP_041636598.1
nicotinamide phosphoribosyltransferase-like	Anguilla anguilla	86.54%	491	XP_035257679.1
nicotinamide phosphoribosyltransferase isoform X1	Loxodonta africana	95.40%	467	XP_023400289.1

nicotinamide phosphoribosyltransferase	Salvelinus alpinus	88.05%	491	XP_023825296.1
nicotinamide phosphoribosyltransferase	Oncorhynchus mykiss	88.05%	491	XP_036813484.1
nicotinamide phosphoribosyltransferase isoform X1	Echeneis naucrates	87.63%	491	XP_029351204.1
nicotinamide phosphoribosyltransferase	Melanotaenia boesemani	87.79%	491	XP_041833125.1
nicotinamide phosphoribosyltransferase	Oncorhynchus mykiss	87.84%	491	XP_021419240.2
nicotinamide phosphoribosyltransferase	Myripristis murdjan	88.21%	491	XP_029901501.1
nicotinamide phosphoribosyltransferase	Hippoglossus hippoglossus	88.56%	491	XP_034434003.1
PREDICTED: nicotinamide phosphoribosyltransferase	Austrofundulus limnaeus	88.00%	491	XP_013888388.1
nicotinamide phosphoribosyltransferase isoform X1	Gopherus evgoodei	88.62%	577	XP_030406654.1
nicotinamide phosphoribosyltransferase	Notolabrus celidotus	87.63%	491	XP_034529362.1
nicotinamide phosphoribosyltransferase	Gadus morhua	87.21%	491	XP_030197079.1
nicotinamide phosphoribosyltransferase	Maylandia zebra	88.00%	491	XP_004560960.1
nicotinamide phosphoribosyltransferase	Polyodon spathula	86.83%	491	XP_041113559.1
nicotinamide phosphoribosyltransferase	Geotrypetes seraphini	88.96%	491	XP_033815171.1
nicotinamide phosphoribosyltransferase	Mastacembelus armatus	87.79%	491	XP_026182158.1
PREDICTED: nicotinamide phosphoribosyltransferase	Lates calcarifer	87.79%	491	XP_018540669.1
nicotinamide phosphoribosyltransferase	Xiphias gladius	88.00%	491	XP_040009472.1

nicotinamide phosphoribosyltransferase	Polyodon spathula	88.19%	491	XP_041110766.1
nicotinamide phosphoribosyltransferase	Protobothrops mucrosquamatus	89.32%	513	XP_015669077.1
nicotinamide phosphoribosyltransferase	Oreochromis aureus	87.79%	491	XP_031582369.1
nicotinamide phosphoribosyltransferase	Astyanax mexicanus	86.75%	489	XP_022522298.1
nicotinamide phosphoribosyltransferase	Oreochromis niloticus	87.79%	491	XP_003450758.3
nicotinamide phosphoribosyltransferase isoform X2	Echeneis naucrates	87.63%	502	XP_029351205.1
nicotinamide phosphoribosyltransferase	Xenopus tropicalis	88.87%	491	NP_001011388.1
nicotinamide phosphoribosyltransferase	Betta splendens	87.18%	490	XP_029018363.1
nicotinamide phosphoribosyltransferase isoform X2	Simochromis diagramma	88.00%	491	XP_039891119.1
nicotinamide phosphoribosyltransferase	Micropterus salmoides	87.58%	491	XP_038594820.1
nicotinamide phosphoribosyltransferase isoform X2	Scophthalmus maximus	86.95%	491	XP_035505590.1
nicotinamide phosphoribosyltransferase	Takifugu rubripes	87.42%	491	XP_003972868.1
nicotinamide phosphoribosyltransferase	Larimichthys crocea	86.04%	491	NP_001290284.1
nicotinamide phosphoribosyltransferase	Chelmon rostratus	87.76%	491	XP_041820227.1
nicotinamide phosphoribosyltransferase	Parambassis ranga	87.16%	491	XP_028280866.1
nicotinamide phosphoribosyltransferase	Periophthalmus magnuspinnatus	87.18%	491	XP_033845046.1
nicotinamide phosphoribosyltransferase	Pelodiscus sinensis	90.27%	484	XP_006137627.2

nicotinamide phosphoribosyltransferase	Rana temporaria	88.75%	498	XP_040199461.1
nicotinamide phosphoribosyltransferase	Morone saxatilis	87.37%	491	XP_035532872.1
nicotinamide phosphoribosyltransferase	Monopterus albus	86.16%	491	XP_020455232.1
PREDICTED: nicotinamide phosphoribosyltransferase	Nanorana parkeri	87.32%	499	XP_018424129.1
nicotinamide phosphoribosyltransferase	Bufo bufo	88.48%	491	XP_040268687.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Dipodomys ordii	87.37%	446	XP_012890189.1
nicotinamide phosphoribosyltransferase isoform X2	Artibeus jamaicensis	87.98%	444	XP_037006387.1
uncharacterized protein LOC735059	Xenopus laevis	88.25%	491	NP_001089988.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Labrus bergylta	86.79%	491	XP_020499699.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Trichosurus vulpecula	86.15%	488	XP_036608359.1
nicotinamide phosphoribosyltransferase	Thalassophryne amazonica	86.11%	491	XP_034018931.1
nicotinamide phosphoribosyltransferase isoform X1	Bos taurus	87.37%	444	XP_010802469.1
nicotinamide phosphoribosyltransferase-like	Salvelinus namaycush	84.71%	489	XP_038873839.1
nicotinamide phosphoribosyltransferase-like	Coregonus clupeaformis	84.71%	489	XP_041692751.1
nicotinamide phosphoribosyltransferase isoform X2	Castor canadensis	87.37%	444	XP_020042027.1
nicotinamide phosphoribosyltransferase isoform X2	Pipistrellus kuhlii	86.56%	444	XP_036285605.1
nicotinamide phosphoribosyltransferase	Colossoma macropomum	88.21%	501	XP_036433674.1

nicotinamide phosphoribosyltransferase isoform X3	Onychostruthus taczanowskii	93.46%	465	XP_041261222.1
PREDICTED: nicotinamide phosphoribosyltransferase	Salmo salar	84.50%	489	XP_013981394.1
nicotinamide phosphoribosyltransferase isoform X4	Onychostruthus taczanowskii	93.67%	465	XP_041261223.1
nicotinamide phosphoribosyltransferase	Lynx canadensis	95.49%	443	XP_030163639.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Gouania willdenowi	84.68%	497	XP_028305847.1
nicotinamide phosphoribosyltransferase isoform X4	Myotis lucifugus	86.35%	444	XP_014308299.1
nicotinamide phosphoribosyltransferase	Clupea harengus	86.60%	491	XP_012679000.2
nicotinamide phosphoribosyltransferase-like isoform X1	Salmo trutta	84.30%	489	XP_029615274.1
nicotinamide phosphoribosyltransferase	Pygocentrus nattereri	86.39%	489	XP_017542057.1
nicotinamide phosphoribosyltransferase	Oncorhynchus mykiss	84.09%	489	XP_021433529.1
nicotinamide phosphoribosyltransferase-like	Oncorhynchus tshawytscha	84.09%	489	XP_024262618.1
nicotinamide phosphoribosyltransferase	Esox lucius	85.06%	489	XP_010901526.1
nicotinamide phosphoribosyltransferase-like	Oncorhynchus kisutch	83.88%	489	XP_020311588.1
nicotinamide phosphoribosyltransferase	Amblyraja radiata	83.78%	488	XP_032893553.1
nicotinamide phosphoribosyltransferase isoform X1	Scophthalmus maximus	85.53%	492	XP_035505580.1
nicotinamide phosphoribosyltransferase-like	Oncorhynchus nerka	84.09%	489	XP_029500570.1
nicotinamide phosphoribosyltransferase isoform X1	Simochromis diagramma	85.95%	500	XP_039891118.1

nicotinamide phosphoribosyltransferase isoform X1	Boleophthalmus pectinirostris	86.04%	490	XP_020782114.1
PREDICTED: nicotinamide phosphoribosyltransferase	Ictalurus punctatus	86.79%	491	XP_017349776.1
PREDICTED: nicotinamide phosphoribosyltransferase	Paralichthys olivaceus	86.75%	492	XP_019947687.1
nicotinamide phosphoribosyltransferase	Cynoglossus semilaevis	85.54%	491	XP_008314007.1
nicotinamide phosphoribosyltransferase isoform X2	Trichechus manatus latirostris	87.58%	451	XP_023592824.1
nicotinamide phosphoribosyltransferase isoform X3	Simochromis diagramma	88.10%	479	XP_039891120.1
nicotinamide phosphoribosyltransferase	Tachysurus fulvidraco	87.21%	491	XP_027006814.1
nicotinamide phosphoribosyltransferase isoform X2	Trachemys scripta elegans	86.46%	497	XP_034622448.1
nicotinamide phosphoribosyltransferase isoform X2	Boleophthalmus pectinirostris	86.04%	489	XP_020782115.1
nicotinamide phosphoribosyltransferase	Pangasianodon hypophthalmus	87.00%	491	XP_026779063.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Cyprinus carpio	88.50%	459	XP_018920679.1
nicotinamide phosphoribosyltransferase	Meleagris gallopavo	95.19%	476	XP_031407977.1
nicotinamide phosphoribosyltransferase	Rhincodon typus	85.77%	490	XP_020371972.1
nicotinamide phosphoribosyltransferase isoform X2	Tursiops truncatus	96.67%	425	XP_019796159.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X2	Ceratotherium simum simum	87.37%	449	XP_014649961.1
nicotinamide phosphoribosyltransferase isoform X2	Equus caballus	86.76%	444	XP_023494645.1
PREDICTED: nicotinamide phosphoribosyltransferase	Callorhinchus milii	82.77%	490	XP_007903337.1

nicotinamide phosphoribosyltransferase-like	Scleropages formosus	86.76%	491	XP_018602971.1
nicotinamide phosphoribosyltransferase	Aotus nancymaae	96.88%	416	XP_021528944.1
nicotinamide phosphoribosyltransferase isoform X2	Manis pentadactyla	84.93%	451	XP_036748387.1
nicotinamide phosphoribosyltransferase isoform X2	Manis javanica	84.73%	451	XP_036859506.1
nicotinamide phosphoribosyltransferase isoform X2	Myotis myotis	86.15%	444	XP_036183439.1
nicotinamide phosphoribosyltransferase isoform X1	Scyliorhinus canicula	84.45%	490	XP_038636087.1
nicotinamide phosphoribosyltransferase isoform X2	Eptesicus fuscus	85.54%	444	XP_028014642.1
nicotinamide phosphoribosyltransferase isoform X1	Carcharodon carcharias	83.89%	490	XP_041071803.1
nicotinamide phosphoribosyltransferase isoform X3	Pipistrellus kuhlii	84.93%	438	XP_036285606.1
nicotinamide phosphoribosyltransferase isoform X2	Octodon degus	96.13%	425	XP_023575280.1
nicotinamide phosphoribosyltransferase isoform X3	Myotis myotis	84.52%	438	XP_036183440.1
nicotinamide phosphoribosyltransferase isoform X2	Delphinapterus leucas	95.53%	469	XP_030617327.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X4	Galeopterus variegatus	96.00%	469	XP_008566098.1
nicotinamide phosphoribosyltransferase isoform X3	Trichechus manatus latirostris	84.93%	438	XP_023592825.1
nicotinamide phosphoribosyltransferase-like	Paramormyrops kingsleyae	85.56%	491	XP_023651821.1
uncharacterized protein LOC557816	Danio rerio	80.50%	1005	XP_021328060.1
nicotinamide phosphoribosyltransferase isoform X2	Meriones unguiculatus	94.82%	430	XP_021501567.1

nicotinamide phosphoribosyltransferase isoform X3	Myotis lucifugus	83.03%	448	XP_023606847.1
uncharacterized protein si:dkey-145c18.2	Danio rerio	80.29%	969	XP_009292044.2
nicotinamide phosphoribosyltransferase isoform X2	Scyliorhinus canicula	83.62%	486	XP_038636088.1
nicotinamide phosphoribosyltransferase	Pantherophis guttatus	83.91%	467	XP_034289565.1
nicotinamide phosphoribosyltransferase isoform X3	Eptesicus fuscus	83.71%	438	XP_028014653.1
nicotinamide phosphoribosyltransferase isoform X3	Scyliorhinus canicula	84.27%	479	XP_038636089.1
PREDICTED: nicotinamide phosphoribosyltransferase isoform X5	Galeopterus variegatus	85.77%	466	XP_008566103.1
nicotinamide phosphoribosyltransferase isoform X3	Manis pentadactyla	92.42%	425	XP_036748388.1
nicotinamide phosphoribosyltransferase isoform X3	Manis javanica	92.42%	425	XP_017535379.1
nicotinamide phosphoribosyltransferase isoform X2	Pan troglodytes	99.74%	404	XP_016801132.1
nicotinamide phosphoribosyltransferase isoform X5	Gopherus evgoodei	89.51%	487	XP_030406990.1
nicotinamide phosphoribosyltransferase	Danio rerio	77.99%	720	XP_021323615.1
nicotinamide phosphoribosyltransferase	Erpetoichthys calabaricus	90.00%	411	XP_028665781.1
nicotinamide phosphoribosyltransferase isoform X4	Manis javanica	92.61%	406	XP_017535387.1
nicotinamide phosphoribosyltransferase isoform X4	Manis pentadactyla	92.61%	406	XP_036748389.1
nicotinamide phosphoribosyltransferase isoform X2	Choloepus didactylus	94.59%	413	XP_037692336.1
nicotinamide phosphoribosyltransferase isoform X6	Myotis lucifugus	96.42%	393	XP_023606849.1

nicotinamide phosphoribosyltransferase isoform X5	Chelonia mydas	93.92%	423	XP_037745080.1
PREDICTED: nicotinamide phosphoribosyltransferase	Eurypyga helias	95.64%	344	XP_010160012.1
nicotinamide phosphoribosyltransferase isoform X2	Carcharodon carcharias	84.70%	403	XP_041071804.1
nicotinamide phosphoribosyltransferase-like	Oncorhynchus keta	87.15%	375	XP_035617204.1
nicotinamide phosphoribosyltransferase-like isoform X1	Petromyzon marinus	61.11%	503	XP_032819019.1
nicotinamide phosphoribosyltransferase isoform X5	Myotis lucifugus	96.60%	404	XP_023606848.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Callorhinchus milii	61.25%	487	XP_007893365.1
PREDICTED: synaptopodin isoform X4	Latimeria chalumnae	61.92%	493	XP_014346999.1
nicotinamide phosphoribosyltransferase 2	Micropterus salmoides	62.16%	493	XP_038552785.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Lepisosteus oculatus	62.13%	497	XP_015202715.1
nicotinamide phosphoribosyltransferase 2	Sander lucioperca	62.16%	490	XP_031133108.1
nicotinamide phosphoribosyltransferase-like	Perca flavescens	62.16%	492	XP_028432105.1
nicotinamide phosphoribosyltransferase 2	Perca fluviatilis	62.16%	492	XP_039652668.1
nicotinamide phosphoribosyltransferase 2	Etheostoma cragini	61.95%	492	XP_034724861.1
nicotinamide phosphoribosyltransferase-like	Sparus aurata	61.54%	578	XP_030276637.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus grahami	61.92%	498	XP_016134261.1
nicotinamide phosphoribosyltransferase 2	Chelmon rostratus	61.95%	493	XP_041808059.1

nicotinamide phosphoribosyltransferase-like	Seriola dumerili	61.75%	493	XP_022600889.1
nicotinamide phosphoribosyltransferase 2	Cheilinus undulatus	61.75%	493	XP_041638749.1
nicotinamide phosphoribosyltransferase-like isoform X2	Chanos chanos	61.07%	500	XP_030633478.1
nicotinamide phosphoribosyltransferase 2	Melanotaenia boesemani	62.16%	493	XP_041836055.1
nicotinamide phosphoribosyltransferase 2	Periophthalmus magnuspinnatus	61.51%	493	XP_033821870.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus rhinoceros	62.34%	498	XP_016406850.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus grahami	62.34%	498	XP_016087284.1
nicotinamide phosphoribosyltransferase-like	Seriola lalandi dorsalis	61.92%	493	XP_023280513.1
nicotinamide phosphoribosyltransferase 2	Epinephelus lanceolatus	62.37%	493	XP_033482800.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus anshuiensis	62.34%	498	XP_016352862.1
nicotinamide phosphoribosyltransferase 2	Morone saxatilis	61.95%	493	XP_035518291.1
nicotinamide phosphoribosyltransferase-like	Echeneis naucrates	61.95%	494	XP_029357812.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Paralichthys olivaceus	61.33%	491	XP_019959881.1
nicotinamide phosphoribosyltransferase 2 isoform X1	Hippoglossus hippoglossus	61.75%	493	XP_034442728.1
nicotinamide phosphoribosyltransferase 2	Toxotes jaculatrix	61.75%	493	XP_040890437.1
nicotinamide phosphoribosyltransferase 2	Hippoglossus stenolepis	61.75%	493	XP_035008120.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Lates calcarifer	61.75%	493	XP_018517282.1

PREDICTED: nicotinamide phosphoribosyltransferase-like	Sinocyclocheilus rhinoceros	61.72%	498	XP_016394343.1
nicotinamide phosphoribosyltransferase-like	Boleophthalmus pectinirostris	61.51%	493	XP_020786623.1
nicotinamide phosphoribosyltransferase-like	Carassius auratus	62.13%	498	XP_026077192.1
nicotinamide phosphoribosyltransferase	Oryzias latipes	61.64%	491	XP_004068957.1
nicotinamide phosphoribosyltransferase-like	Amblyraja radiata	61.04%	473	XP_032892535.1
nicotinamide phosphoribosyltransferase-like	Myripristis murdjan	61.07%	493	XP_029908295.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase 2	Scophthalmus maximus	61.27%	493	XP_035487220.1
nicotinamide phosphoribosyltransferase 2	Amphiprion ocellaris	61.33%	493	XP_023138323.1
nicotinamide phosphoribosyltransferase 2	Salvelinus namaycush	61.17%	498	XP_038864904.1
nicotinamide phosphoribosyltransferase-like	Parambassis ranga	61.92%	487	XP_028262149.1
nicotinamide phosphoribosyltransferase	Larimichthys crocea	61.75%	492	XP_010733618.1
nicotinamide phosphoribosyltransferase 2	Oryzias melastigma	60.96%	491	XP_024126447.1
nicotinamide phosphoribosyltransferase 2	Megalops cyprinoides	61.72%	497	XP_036387526.1
nicotinamide phosphoribosyltransferase-like isoform X1	Petromyzon marinus	59.50%	491	XP_032827368.1
nicotinamide phosphoribosyltransferase 2	Kryptolebias marmoratus	61.83%	496	XP_017279238.1
nicotinamide phosphoribosyltransferase 2 isoform X1	Xiphias gladius	61.54%	493	XP_040015386.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Hippocampus comes	61.03%	532	XP_019723247.1

nicotinamide phosphoribosyltransferase 2	Colossoma macropomum	60.67%	497	XP_036415034.1
nicotinamide phosphoribosyltransferase 2	Pimephales promelas	61.09%	498	XP_039529555.1
nicotinamide phosphoribosyltransferase-like isoform X1	Chanos chanos	61.07%	497	XP_030633477.1
nicotinamide phosphoribosyltransferase 2	Esox lucius	60.33%	498	XP_010879972.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X2	Poecilia latipinna	61.92%	493	XP_014885998.1
nicotinamide phosphoribosyltransferase-like isoform X2	Paramormyrops kingsleyae	61.09%	494	XP_023649794.1
nicotinamide phosphoribosyltransferase 2	Carcharodon carcharias	59.17%	493	XP_041047986.1
nicotinamide phosphoribosyltransferase 2	Pangasianodon hypophthalmus	59.84%	497	XP_026791198.1
nicotinamide phosphoribosyltransferase 2	Anguilla anguilla	61.09%	497	XP_035244758.1
nicotinamide phosphoribosyltransferase-like isoform X1	Xiphophorus couchianus	61.51%	493	XP_027859448.1
nicotinamide phosphoribosyltransferase 2	Scyliorhinus canicula	59.50%	502	XP_038666491.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Salmo salar	60.75%	498	XP_013988920.1
nicotinamide phosphoribosyltransferase-like isoform X3	Polyodon spathula	60.88%	497	XP_041080406.1
nicotinamide phosphoribosyltransferase-like	Acanthochromis polyacanthus	60.91%	493	XP_022075469.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X2	Poecilia mexicana	61.59%	493	XP_014866798.1
nicotinamide phosphoribosyltransferase-like	Astyanax mexicanus	60.25%	497	XP_022538522.1
nicotinamide phosphoribosyltransferase-like isoform X1	Xiphophorus maculatus	61.51%	493	XP_005803755.1

nicotinamide phosphoribosyltransferase-like	Salmo trutta	60.75%	498	XP_029630853.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X2	Poecilia formosa	61.59%	493	XP_007569917.1
nicotinamide phosphoribosyltransferase-like isoform X1	Xiphophorus hellerii	61.72%	493	XP_032405949.1
nicotinamide phosphoribosyltransferase 2	Sebastes umbrosus	61.33%	493	XP_037620968.1
nicotinamide phosphoribosyltransferase 2	Pungitius pungitius	60.91%	493	XP_037346824.1
nicotinamide phosphoribosyltransferase-like	Sphaeramia orbicularis	60.75%	493	XP_029989506.1
nicotinamide phosphoribosyltransferase 2	Danio rerio	61.12%	498	NP_997833.2
nicotinamide phosphoribosyltransferase-like	Carassius auratus	61.30%	498	XP_026104562.1
nicotinamide phosphoribosyltransferase 2	Cyprinodon tularosa	61.54%	493	XP_038124996.1
nicotinamide phosphoribosyltransferase 2 isoform X1	Betta splendens	61.51%	492	XP_029006650.1
nicotinamide phosphoribosyltransferase 2	Polypterus senegalus	60.67%	494	XP_039626513.1
nicotinamide phosphoribosyltransferase 2	Electrophorus electricus	60.25%	497	XP_035376379.1
nicotinamide phosphoribosyltransferase-like isoform X2	Oncorhynchus nerka	60.33%	498	XP_029539442.1
nicotinamide phosphoribosyltransferase 2	Notolabrus celidotus	60.67%	493	XP_034546750.1
nicotinamide phosphoribosyltransferase 2 isoform X1	Oncorhynchus mykiss	60.46%	498	XP_021425277.2
nicotinamide phosphoribosyltransferase 2 isoform X1	Oncorhynchus keta	60.33%	498	XP_035633411.1
nicotinamide phosphoribosyltransferase isoform X1	Oncorhynchus kisutch	60.46%	498	XP_020347048.1

nicotinamide phosphoribosyltransferase-like isoform X1	Oncorhynchus tshawytscha	60.46%	498	XP_024236080.1
nicotinamide phosphoribosyltransferase 2	Cyclopterus lumpus	60.91%	493	XP_034388595.1
nicotinamide phosphoribosyltransferase-like	Acipenser ruthenus	60.46%	497	XP_034776163.1
nicotinamide phosphoribosyltransferase-like	Acipenser ruthenus	60.25%	497	XP_033879900.1
nicotinamide phosphoribosyltransferase 2	Pygocentrus nattereri	60.46%	497	XP_017567241.1
nicotinamide phosphoribosyltransferase-like	Tachysurus fulvidraco	59.75%	497	XP_027032127.1
nicotinamide phosphoribosyltransferase-like isoform X1	Monopterus albus	60.50%	493	XP_020441598.1
nicotinamide phosphoribosyltransferase-like	Scleropages formosus	61.09%	498	XP_018589722.1
nicotinamide phosphoribosyltransferase-like	Erpetoichthys calabaricus	60.25%	494	XP_028679932.1
nicotinamide phosphoribosyltransferase-like	Denticeps clupeoides	60.25%	497	XP_028855317.1
nicotinamide phosphoribosyltransferase-like	Clupea harengus	59.46%	497	XP_012696704.2
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X1	Poecilia latipinna	59.56%	511	XP_014885997.1
nicotinamide phosphoribosyltransferase-like isoform X2	Gadus morhua	59.59%	496	XP_030230559.1
PREDICTED: synaptoporin isoform X3	Latimeria chalumnae	61.56%	505	XP_014346997.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X1	Poecilia mexicana	59.24%	513	XP_014866797.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase 2	Coregonus clupeaformis	60.33%	496	XP_041743587.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X1	Poecilia formosa	59.24%	513	XP_007569916.1

nicotinamide phosphoribosyltransferase-like	Polyodon spathula	59.62%	492	XP_041130896.1
nicotinamide phosphoribosyltransferase-like	Branchiostoma floridae	59.50%	502	XP_035677089.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X1	Ictalurus punctatus	58.81%	497	XP_017306101.1
nicotinamide phosphoribosyltransferase-like	Anarrhichthys ocellatus	59.42%	501	XP_031731460.1
nicotinamide phosphoribosyltransferase 2	Acanthopagrus latus	60.47%	488	XP_036957919.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Austrofundulus limnaeus	59.83%	494	XP_013881832.1
nicotinamide phosphoribosyltransferase-like isoform X2	Polyodon spathula	60.30%	507	XP_041080405.1
nicotinamide phosphoribosyltransferase	Lingula anatina	60.17%	482	XP_013415217.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Branchiostoma belcheri	59.29%	482	XP_019633331.1
nicotinamide phosphoribosyltransferase-like isoform X2	Cottoperca gobio	61.12%	493	XP_029288169.1
nicotinamide phosphoribosyltransferase-like isoform X1	Polyodon spathula	56.73%	532	XP_041080404.1
nicotinamide phosphoribosyltransferase 2 isoform X1	Gasterosteus aculeatus aculeatus	60.91%	493	XP_040059081.1
nicotinamide phosphoribosyltransferase	Exaiptasia diaphana	58.99%	480	XP_020913725.1
nicotinamide phosphoribosyltransferase-like	Gouania willdenowi	59.46%	491	XP_028302967.1
PREDICTED: nicotinamide phosphoribosyltransferase	Balearica regulorum gibbericeps	93.67%	337	XP_010296937.1
nicotinamide phosphoribosyltransferase	Nematostella vectensis	58.08%	475	XP_032226992.1
nicotinamide phosphoribosyltransferase-like isoform X1	Paramormyrops kingsleyae	60.00%	512	XP_023649793.1

nicotinamide phosphoribosyltransferase-like	Etheostoma spectabile	62.50%	446	XP_032369597.1
nicotinamide phosphoribosyltransferase-like	Crassostrea virginica	60.00%	469	XP_022301606.1
nicotinamide phosphoribosyltransferase-like	Pomacea canaliculata	60.04%	477	XP_025106706.1
nicotinamide phosphoribosyltransferase 2	Pseudochaenichthys georgianus	59.88%	493	XP_033938110.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Stegastes partitus	60.09%	477	XP_008303670.1
nicotinamide phosphoribosyltransferase 2	Gymnodraco acuticeps	59.67%	493	XP_034089288.1
nicotinamide phosphoribosyltransferase 2	Trematomus bernacchii	59.46%	493	XP_033999684.1
nicotinamide phosphoribosyltransferase-like isoform X1	Oncorhynchus nerka	59.70%	559	XP_029539441.1
nicotinamide phosphoribosyltransferase-like	Gigantopelta aegis	58.18%	478	XP_041347475.1
PREDICTED: nicotinamide phosphoribosyltransferase	Mesitornis unicolor	95.39%	304	XP_010187927.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X2	Ictalurus punctatus	58.65%	486	XP_017306102.1
nicotinamide phosphoribosyltransferase-like isoform X1	Gadus morhua	59.15%	505	XP_030230558.1
nicotinamide phosphoribosyltransferase-like	Mizuhopecten yessoensis	57.35%	481	XP_021368938.1
nicotinamide phosphoribosyltransferase	Crassostrea gigas	58.74%	469	XP_011447799.2
PREDICTED: nicotinamide phosphoribosyltransferase-like	Amphimedon queenslandica	59.21%	466	XP_003385030.1
nicotinamide phosphoribosyltransferase	Aplysia californica	56.63%	481	XP_012942330.1
nicotinamide phosphoribosyltransferase-like	Pecten maximus	58.82%	484	XP_033744427.1

nicotinamide phosphoribosyltransferase-like isoform X1	Cottoperca gobio	59.18%	520	XP_029288168.1
nicotinamide phosphoribosyltransferase-like isoform X1	Pocillopora damicornis	57.80%	481	XP_027040863.1
nicotinamide phosphoribosyltransferase-like isoform X4	Cottoperca gobio	60.73%	478	XP_029288171.1
nicotinamide phosphoribosyltransferase-like isoform X3	Cottoperca gobio	60.73%	489	XP_029288170.1
nicotinamide phosphoribosyltransferase-like	Stylophora pistillata	56.76%	481	XP_022790158.1
nicotinamide phosphoribosyltransferase-like	Octopus sinensis	57.77%	474	XP_036357641.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Octopus bimaculoides	57.77%	474	XP_014783337.1
nicotinamide phosphoribosyltransferase-like	Actinia tenebrosa	55.95%	475	XP_031574499.1
pre-B-cell colony-enhancing factor	Capsaspora owczarzaki ATCC 30864	57.14%	469	XP_004363953.2
nicotinamide phosphoribosyltransferase-like	Strongylocentrotus purpuratus	57.56%	480	XP_030829364.1
nicotinamide phosphoribosyltransferase	Strongylocentrotus purpuratus	57.35%	480	XP_782393.1
nicotinamide phosphoribosyltransferase-like	Lytechinus variegatus	57.14%	480	XP_041460637.1
nicotinamide phosphoribosyltransferase-like isoform X2	Pocillopora damicornis	57.20%	471	XP_027040864.1
Quinolate phosphoribosyl transferase,C-terminal domain and Nicotinamide phosphoribosyl	Strongyloides ratti	56.96%	506	XP_024498824.1
nicotinamide phosphoribosyltransferase-like	Anneissia japonica	58.40%	472	XP_033099752.1
nicotinamide phosphoribosyltransferase-like isoform X1	Limulus polyphemus	56.30%	475	XP_022253723.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Actinia tenebrosa	55.53%	491	XP_031567082.1

nicotinamide phosphoribosyltransferase-like	Anneissia japonica	58.19%	472	XP_033104415.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Biomphalaria glabrata	57.82%	458	XP_013090296.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Saccoglossus kowalevskii	58.53%	478	XP_002741603.1
nicotinamide phosphoribosyltransferase-like	Acropora millepora	56.21%	480	XP_029196192.1
uncharacterized protein MONBRDRAFT_24677	Monosiga brevicollis MX1	54.95%	508	XP_001744923.1
nicotinamide phosphoribosyltransferase isoform X2	Octopus sinensis	57.05%	478	XP_036358017.1
nicotinamide phosphoribosyltransferase-like	Acanthaster planci	57.49%	475	XP_022096002.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Octopus bimaculoides	56.63%	478	XP_014789529.1
hypothetical protein TRIADDRAFT_20412	Trichoplax adhaerens	56.12%	469	XP_002109387.1
nicotinamide phosphoribosyltransferase-like	Asterias rubens	58.00%	475	XP_033647974.1
nicotinamide phosphoribosyltransferase-like isoform X1	Patiria miniata	56.79%	487	XP_038061094.1
nicotinamide phosphoribosyltransferase-like isoform X2	Patiria miniata	56.79%	472	XP_038061095.1
nicotinamide phosphoribosyltransferase-like	Limulus polyphemus	55.49%	475	XP_013785620.1
Nicotinamide phosphoribosyltransferase	Echinococcus granulosus	57.14%	589	XP_024349285.1
nicotinamide phosphoribosyltransferase-like isoform X1	Penaeus monodon	56.93%	485	XP_037798587.1
nicotinamide phosphoribosyltransferase-like	Penaeus vannamei	56.81%	487	XP_027236932.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Amphimedon queenslandica	54.30%	480	XP_003384858.1

nicotinamide phosphoribosyltransferase isoform X1	Octopus sinensis	56.65%	460	XP_036358016.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Elephantulus edwardii	95.96%	291	XP_006894663.1
nicotinamide phosphoribosyltransferase 2 isoform X1	Anabas testudineus	61.27%	406	XP_026200359.1
nicotinamide phosphoribosyltransferase-like isoform X3	Petromyzon marinus	60.49%	427	XP_032819021.1
Pre-B cell enhancing factor precursor	Brugia malayi	55.13%	498	XP_001893248.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Priapulus caudatus	53.68%	475	XP_014673125.1
nicotinamide phosphoribosyltransferase 2 isoform X2	Betta splendens	55.86%	445	XP_029006651.1
pre-B cell enhancing factor	Loa loa	53.78%	494	XP_003144417.1
LOW QUALITY PROTEIN: nicotinamide phosphoribosyltransferase-like	Manis pentadactyla	82.89%	342	XP_036749481.1
hypothetical protein T265_00471	Opisthorchis viverrini	52.27%	515	XP_009162508.1
nicotinamide phosphoribosyltransferase 2 isoform X2	Hippoglossus hippoglossus	61.87%	415	XP_034442729.1
nicotinamide phosphoribosyltransferase-like isoform X2	Monopterus albus	53.22%	450	XP_020441599.1
hypothetical protein LOTGIDRAFT_218343	Lottia gigantea	53.88%	469	XP_009059352.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X3	Poecilia latipinna	61.83%	415	XP_014885999.1
nicotinamide phosphoribosyltransferase-like	Spodoptera litura	53.26%	457	XP_022836031.1
nicotinamide phosphoribosyltransferase-like isoform X2	Xiphophorus couchianus	61.32%	415	XP_027859449.1
nicotinamide phosphoribosyltransferase-like isoform X2	Xiphophorus hellerii	61.58%	415	XP_032405950.1

PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X2	Poecilia reticulata	61.58%	415	XP_008408569.1
nicotinamide phosphoribosyltransferase-like isoform X2	Xiphophorus maculatus	61.32%	415	XP_023180742.1
nicotinamide phosphoribosyltransferase 2 isoform X2	Gasterosteus aculeatus aculeatus	60.35%	415	XP_040059083.1
nicotinate phosphoribosyltransferase	Arcicella rosea	52.84%	457	WP_184134646.1
nicotinate phosphoribosyltransferase, (NAPRTase) family	Necator americanus	52.42%	472	XP_013301963.1
nicotinamide phosphoribosyltransferase-like isoform X2	Oncorhynchus tshawytscha	59.80%	419	XP_024236087.1
nicotinamide phosphoribosyltransferase isoform X2	Oncorhynchus kisutch	59.80%	419	XP_031662171.1
nicotinamide phosphoribosyltransferase 2 isoform X2	Oncorhynchus mykiss	59.80%	419	XP_036806453.1
nicotinamide phosphoribosyltransferase-like isoform X3	Oncorhynchus nerka	59.39%	419	XP_029539443.1
nicotinate phosphoribosyltransferase	Pseudarcicella hirudinis	52.74%	459	WP_092013600.1
nicotinate phosphoribosyltransferase related pre-B cell enhancing factor	Schistosoma mansoni	53.09%	472	XP_018650645.1
nicotinamide phosphoribosyltransferase 2 isoform X2	Oncorhynchus keta	59.39%	419	XP_035633412.1
PREDICTED: nicotinamide phosphoribosyltransferase-like isoform X3	Ictalurus punctatus	58.81%	419	XP_017306103.1
nicotinate phosphoribosyltransferase	Flectobacillus sp. BAB-3569	53.39%	456	WP_095161636.1
nicotinate phosphoribosyltransferase	Arcicella aurantiaca	51.67%	467	WP_109743025.1
nicotinate phosphoribosyltransferase	Cellulophaga sp. BC115SP	53.39%	456	WP_166550341.1
nicotinate phosphoribosyltransferase	Flectobacillus major	52.53%	457	WP_026995933.1

nicotinate phosphoribosyltransferase	Emticicia sp. ODNR4P	52.75%	456	WP_166579510.1
nicotinamide phosphoribosyltransferase-like isoform X5	Cottoperca gobio	60.61%	415	XP_029288173.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Stegastes partitus	62.67%	351	XP_008290401.1
hypothetical protein HELRODRAFT_67798	Helobdella robusta	52.80%	456	XP_009025138.1
hypothetical protein AURANDRAFT_29036	Aureococcus anophagefferens	51.69%	457	XP_009038694.1
nicotinate phosphoribosyltransferase	Sandaracinus amylolyticus	48.73%	460	WP_053238714.1
nicotinate phosphoribosyltransferase	Gloeocapsa sp. PCC 73106	51.05%	461	WP_006529042.1
hypothetical protein AURANDRAFT_39495	Aureococcus anophagefferens	51.69%	497	XP_009042743.1
nicotinamide phosphoribosyltransferase-like isoform X2	Penaeus monodon	58.52%	419	XP_037798589.1
nicotinamide phosphoribosyltransferase-like	Eurytemora affinis	51.27%	466	XP_023333671.1
nicotinate phosphoribosyltransferase	Fimbrioglobus ruber	50.64%	471	WP_202974134.1
nicotinate phosphoribosyltransferase	Pseudobacteriovorax antillogorgiicola	50.84%	464	WP_132316807.1
nicotinamide phosphoribosyltransferase-like	Penaeus monodon	49.89%	481	XP_037798270.1
nicotinamide phosphoribosyltransferase-like	Penaeus vannamei	48.63%	481	XP_027212282.1
nicotinate phosphoribosyltransferase	Chitinophaga terrae Kim and Jung 2007	49.17%	471	WP_089765143.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. GDMCC 1.1288	48.15%	471	WP_116975716.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. 1303	49.17%	471	WP_168861791.1

nicotinate phosphoribosyltransferase	Chitinophaga eiseniae	49.17%	471	WP_078669551.1
nicotinate phosphoribosyltransferase	Chitinophaga fulva	49.07%	471	WP_169223454.1
nicotinate phosphoribosyltransferase	Chitinophaga arvensicola	47.73%	471	WP_089890861.1
nicotinate phosphoribosyltransferase	Microbulbifer thermotolerans	47.92%	473	WP_074901071.1
nicotinate phosphoribosyltransferase	Microbulbifer thermotolerans	47.92%	473	WP_067153313.1
nicotinate phosphoribosyltransferase	Chitinophaga ginsengisegetis	48.45%	471	WP_079471630.1
nicotinate phosphoribosyltransferase	Oleiphilus messinensis	47.48%	469	WP_087462700.1
nicotinate phosphoribosyltransferase	Thalassocella blandensis	48.10%	463	WP_142594781.1
nicotinate phosphoribosyltransferase	Chitinophaga niastensis	47.93%	471	WP_106526668.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. KRA15-503	48.76%	471	WP_212001120.1
nicotinate phosphoribosyltransferase	Chitinophaga varians	47.84%	471	WP_188097204.1
PREDICTED: synaptoporin isoform X2	Latimeria chalumnae	64.00%	580	XP_014346996.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Poecilia mexicana	84.49%	263	XP_014825798.1
nicotinate phosphoribosyltransferase	Chitinophaga caeni	47.94%	473	WP_098192935.1
nicotinate phosphoribosyltransferase	Pseudoflavitalea sp. G-6-1-2	48.06%	481	WP_169424074.1
nicotinate phosphoribosyltransferase	Chitinophaga flava	48.55%	471	WP_113615967.1
PREDICTED: nicotinamide phosphoribosyltransferase-like	Poecilia formosa	84.08%	263	XP_007540734.1

nicotinate phosphoribosyltransferase	Chitinophaga varians	47.94%	471	WP_168870727.1
nicotinate phosphoribosyltransferase	Chitinophaga eiseniae	47.93%	469	WP_168738414.1
nicotinate phosphoribosyltransferase	Fulvivirga kasyanovii	47.33%	480	WP_155175453.1
nicotinate phosphoribosyltransferase	Chitinophaga qingshengii	48.14%	471	WP_188088545.1
nicotinate phosphoribosyltransferase	Solitalea canadensis	48.13%	469	WP_014682330.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. CF418	48.35%	471	WP_149694217.1
nicotinate phosphoribosyltransferase	Chitinophaga polysaccharea	47.31%	471	WP_142684411.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. OAE865	47.31%	471	WP_209855631.1
nicotinate phosphoribosyltransferase	Thalassospira sp. MCCC 1A01428	48.42%	458	WP_085589522.1
nicotinate phosphoribosyltransferase	Chitinophaga barathri	47.73%	469	WP_120518914.1
nicotinate phosphoribosyltransferase	Microbulbifer rhizosphaerae	47.49%	471	WP_183457077.1
nicotinate phosphoribosyltransferase	Microbulbifer sp. YPW1	48.02%	471	WP_181253464.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. OK723	47.31%	471	WP_133807103.1
nicotinate phosphoribosyltransferase	Thalassospira mesophila	48.42%	474	WP_085585482.1
nicotinate phosphoribosyltransferase	Microbulbifer taiwanensis	47.50%	460	WP_193190327.1
nicotinate phosphoribosyltransferase	Fulvivirga sp. 29W222	47.63%	480	WP_202856492.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. 2R12	47.73%	471	WP_211971144.1

nicotinate phosphoribosyltransferase	Hahella chejuensis	48.10%	459	WP_011400657.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. SYP-B3965	47.31%	469	WP_153660752.1
nicotinate phosphoribosyltransferase	Niastella populi	47.10%	479	WP_081165586.1
nicotinate phosphoribosyltransferase	Chitinophaga extrema	47.42%	471	WP_127034736.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. S165	48.56%	471	WP_110053546.1
nicotinate phosphoribosyltransferase	Chitinophaga dinghuensis	47.93%	471	WP_111591019.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. O9	47.42%	471	WP_160712245.1
nicotinate phosphoribosyltransferase	Chitinophaga polysaccharea	47.52%	469	WP_145662406.1
nicotinate phosphoribosyltransferase	Hahella sp. KA22	47.89%	459	WP_127974303.1
nicotinamide phosphoribosyltransferase-like	Anarrhichthys ocellatus	86.38%	408	XP_031721473.1
nicotinate phosphoribosyltransferase	Chitinophaga ginsengisoli	47.74%	471	WP_106599999.1
nicotinate phosphoribosyltransferase	Niastella sp. MAH-29	46.83%	480	WP_209137387.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. 1310	49.05%	515	WP_168808747.1
nicotinate phosphoribosyltransferase	Microbulbifer sp. NBU-8HK146	46.23%	460	WP_193163680.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. OAE873	47.63%	471	WP_199654209.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Oleiphilus	47.58%	472	WP_068481096.1
nicotinate phosphoribosyltransferase	Thalassospira marina	47.27%	472	WP_101264106.1

nicotinate phosphoribosyltransferase	Oleibacter marinus	47.68%	458	WP_076514536.1
nicotinate phosphoribosyltransferase	Marivirga lumbricoides	47.96%	476	WP_188463711.1
nicotinate phosphoribosyltransferase	Thalassospira marina	47.16%	472	WP_101284186.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. MAH-28	46.69%	467	WP_209145892.1
nicotinate phosphoribosyltransferase	Chitinophaga niabensis	46.47%	469	WP_074241636.1
nicotinate phosphoribosyltransferase	Chitinophaga vietnamensis	46.69%	471	WP_143309514.1
nicotinate phosphoribosyltransferase	Microbulbifer donghaiensis	47.07%	460	WP_073274293.1
nicotinate phosphoribosyltransferase	Chitinophaga filiformis	48.14%	471	WP_089829265.1
nicotinate phosphoribosyltransferase	Chitinophaga sancti	47.74%	471	WP_183289454.1
nicotinate phosphoribosyltransferase	Chitinophaga sancti	47.74%	471	WP_204549677.1
nicotinate phosphoribosyltransferase	Chitinophaga pinensis	48.35%	471	WP_146302716.1
nicotinate phosphoribosyltransferase	Niastella yeongjuensis	46.68%	480	WP_081200922.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Chitinophaga	47.11%	469	WP_168764855.1
nicotinate phosphoribosyltransferase	Chitinophaga rupis	47.31%	469	WP_089906823.1
nicotinate phosphoribosyltransferase	Paraflavitalea soli	46.47%	481	WP_119052589.1
nicotinate phosphoribosyltransferase	Mucilaginibacter paludis	45.55%	472	WP_008504805.1
nicotinate phosphoribosyltransferase	[Flexibacter] sp. ATCC 35208	47.42%	471	WP_083722002.1

nicotinate phosphoribosyltransferase	Chitinophaga silvisoli	47.53%	471	WP_116852935.1
nicotinamide phosphoribosyltransferase	Ixodes scapularis	46.12%	487	XP_040066246.1
nicotinamide phosphoribosyltransferase 2 isoform X2	Anabas testudineus	60.68%	328	XP_026200360.1
nicotinate phosphoribosyltransferase	Marivirga tractuosa	47.73%	488	WP_013454807.1
nicotinate phosphoribosyltransferase	Chitinophaga agri	47.63%	471	WP_162333214.1
nicotinate phosphoribosyltransferase	Thalassospira sp. TSL5-1	47.16%	471	WP_143181925.1
nicotinate phosphoribosyltransferase	Microbulbifer sp. A4B17	47.70%	470	WP_108731290.1
nicotinate phosphoribosyltransferase	Niaistella koreensis	46.68%	479	WP_014222021.1
nicotinate phosphoribosyltransferase	Variovorax boronicumulans	46.25%	481	WP_062480688.1
nicotinate phosphoribosyltransferase	Chitinophaga pinensis	47.42%	471	WP_012788281.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. YR627	47.42%	471	WP_089810239.1
nicotinate phosphoribosyltransferase	Variovorax boronicumulans	46.25%	481	WP_095950297.1
nicotinate phosphoribosyltransferase	Niaistella caeni	46.42%	479	WP_136578778.1
nicotinate phosphoribosyltransferase	Chitinophaga jiangningensis	46.28%	471	WP_073082267.1
nicotinate phosphoribosyltransferase	Thalassospira profundimaris	46.85%	469	WP_114090309.1
nicotinate phosphoribosyltransferase	Variovorax boronicumulans	46.25%	481	WP_070063034.1
nicotinate phosphoribosyltransferase	Variovorax sp. KBS0712	46.25%	481	WP_139700490.1

nicotinate phosphoribosyltransferase	Amphritea spongicola	47.49%	466	WP_205657313.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Variovorax	45.83%	481	WP_125952314.1
nicotinate phosphoribosyltransferase	Variovorax sp. BK460	45.93%	481	WP_130428394.1
nicotinate phosphoribosyltransferase	Variovorax sp. RO1	46.44%	481	WP_101491400.1
nicotinate phosphoribosyltransferase	Chitinophaga sancti	46.91%	471	WP_072358755.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. ASV3	46.69%	471	WP_199075945.1
nicotinate phosphoribosyltransferase	Chitinophaga skermanii	46.30%	471	WP_111599176.1
nicotinate phosphoribosyltransferase	Variovorax sp. YR750	46.46%	481	WP_093208281.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.57%	458	WP_211033260.1
nicotinate phosphoribosyltransferase	Niastella vici	45.62%	479	WP_081145804.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.99%	458	WP_196762203.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. Cy-1792	46.71%	471	WP_167014376.1
nicotinate phosphoribosyltransferase	Mucilaginibacter lappiensis	45.45%	470	WP_183586648.1
nicotinate phosphoribosyltransferase	Variovorax sp. 38R	46.44%	481	WP_192193982.1
PREDICTED: synaptoporin isoform X5	Latimeria chalumnae	63.55%	439	XP_014347000.1
nicotinate phosphoribosyltransferase	Sporocytophaga myxococcoides	45.04%	485	WP_037576589.1
nicotinate phosphoribosyltransferase	Mucilaginibacter lappiensis	45.45%	470	WP_076372304.1

nicotinate phosphoribosyltransferase	Hahella sp. CCB-MM4	47.48%	477	WP_094706123.1
nicotinamide phosphoribosyltransferase-like	Rhipicephalus microplus	43.53%	521	XP_037280693.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.57%	457	WP_142049914.1
nicotinate phosphoribosyltransferase	Microbulbifer sp. A4B17	46.64%	471	WP_108734716.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.57%	457	WP_063365379.1
nicotinate phosphoribosyltransferase	Microbulbifer marinus	46.44%	460	WP_091388251.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.57%	458	WP_211270614.1
nicotinate phosphoribosyltransferase	Chitinophaga cymbidii	46.50%	469	WP_146859086.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.57%	458	WP_196765657.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Variovorax	46.44%	481	WP_140840795.1
nicotinate phosphoribosyltransferase	Microbulbifer rhizosphaerae	46.35%	460	WP_183459191.1
nicotinate phosphoribosyltransferase	Variovorax sp. SG533	46.25%	484	WP_176661790.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.15%	457	WP_065792041.1
nicotinate phosphoribosyltransferase	Ketobacter sp. MCCC 1A13808	45.53%	471	WP_156873516.1
nicotinate phosphoribosyltransferase	Microbulbifer sp. THAF38	45.82%	460	WP_152450442.1
nicotinate phosphoribosyltransferase	Methylomagnum ishizawai	47.03%	467	WP_085215568.1
nicotinate phosphoribosyltransferase	Variovorax sp. KBW07	46.15%	481	WP_124466142.1

nicotinate phosphoribosyltransferase	Niastella sp. CF465	46.67%	479	WP_132554669.1
nicotinate phosphoribosyltransferase	Chryseolinea sp. Jin1	47.01%	477	WP_202009727.1
nicotinate phosphoribosyltransferase	Alcanivorax profundus	45.70%	467	WP_119917872.1
nicotinate phosphoribosyltransferase	Microbulbifer taiwanensis	45.49%	458	WP_193193860.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.57%	458	WP_196761461.1
nicotinate phosphoribosyltransferase	Vibrio sonorensis	47.25%	455	WP_070965424.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Variovorax	46.04%	481	WP_093058646.1
nicotinate phosphoribosyltransferase	Pseudobacter ginsenosidimutans	47.11%	473	WP_130541977.1
nicotinate phosphoribosyltransferase	Mucilagibacter sp. R11	45.96%	470	WP_166584853.1
nicotinate phosphoribosyltransferase	Variovorax sp. IB41	45.93%	481	WP_198788745.1
nicotinate phosphoribosyltransferase	Microbulbifer sp. GL-2	46.03%	460	WP_143729829.1
nicotinate phosphoribosyltransferase	Xenophilus sp. E41	46.04%	479	WP_121695488.1
nicotinate phosphoribosyltransferase	Chitinophaga oryzae	47.01%	471	WP_157301183.1
nicotinate phosphoribosyltransferase	Variovorax paradoxus	45.93%	485	WP_013544029.1
nicotinate phosphoribosyltransferase	Variovorax sp. 770b2	45.72%	481	WP_093440616.1
nicotinate phosphoribosyltransferase	Variovorax guangxiensis	46.06%	481	WP_126025437.1
nicotinate phosphoribosyltransferase	Variovorax paradoxus	45.93%	481	WP_042581347.1

nicotinate phosphoribosyltransferase	Variovorax guangxiensis	46.06%	481	WP_184639652.1
nicotinate phosphoribosyltransferase	Hahella ganhwensis	46.20%	476	WP_020407480.1
nicotinate phosphoribosyltransferase	Variovorax sp. SG517	45.83%	481	WP_176665082.1
nicotinate phosphoribosyltransferase	Ohtaekwangia koreensis	46.80%	478	WP_079686492.1
nicotinate phosphoribosyltransferase	Marivirga sp. S37H4	47.52%	478	WP_201431452.1
nicotinate phosphoribosyltransferase	Alcanivorax sp. P2S70	45.68%	467	WP_022985890.1
nicotinate phosphoribosyltransferase	Variovorax paradoxus	46.04%	481	WP_019657723.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. OK268	45.85%	470	WP_090466428.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. SG564	45.34%	470	WP_172336552.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Microbulbifer	45.19%	460	WP_067084642.1
nicotinamide phosphoribosyltransferase-like	Rhipicephalus sanguineus	43.30%	547	XP_037521025.1
nicotinate phosphoribosyltransferase	Chitinophaga lutea	46.71%	470	WP_123846113.1
PREDICTED: synaptoporin isoform X1	Latimeria chalumnae	63.55%	592	XP_014346995.1
nicotinate phosphoribosyltransferase	Candidatus Jidaibacter acanthamoeba	48.85%	461	WP_039457084.1
nicotinate phosphoribosyltransferase	Variovorax sp. YR266	46.04%	481	WP_093179760.1
nicotinate phosphoribosyltransferase	Pseudoflavitalea rhizosphaerae	46.90%	473	WP_127128919.1
nicotinate phosphoribosyltransferase	Variovorax sp. 1180	46.04%	481	WP_145759660.1

nicotinate phosphoribosyltransferase	Variovorax sp. OK605	45.72%	481	WP_093133565.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Variovorax	45.72%	481	WP_093219367.1
nicotinate phosphoribosyltransferase	Chitinophaga pinensis	48.56%	471	WP_146968577.1
nicotinate phosphoribosyltransferase	Variovorax sp. BK151	46.04%	481	WP_132756286.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Variovorax	46.04%	481	WP_132802773.1
nicotinate phosphoribosyltransferase	Chroococidiopsis sp. PCC 6712	47.41%	469	WP_169242574.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. ysch24	46.50%	471	WP_157304377.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. YR573	46.79%	471	WP_089781395.1
nicotinate phosphoribosyltransferase	Variovorax sp. Root411	45.51%	481	WP_056516590.1
nicotinate phosphoribosyltransferase	Microbulbifer yueqingensis	45.61%	464	WP_091514300.1
nicotinate phosphoribosyltransferase	Variovorax sp. 369	46.04%	481	WP_126183816.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.89%	458	WP_200890385.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Variovorax	45.72%	481	WP_081267491.1
nicotinate phosphoribosyltransferase	Xenophilus azovorans	45.44%	480	WP_038216004.1
nicotinate phosphoribosyltransferase	Chitinophaga japonensis	45.66%	473	WP_211366105.1
nicotinate phosphoribosyltransferase	Mucilaginibacter inviolabilis	45.85%	470	WP_166088749.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	46.01%	457	WP_200768090.1

nicotinate phosphoribosyltransferase	Xylophilus ampelinus	46.32%	479	WP_066117069.1
nicotinate phosphoribosyltransferase	Mucilaginibacter yixingensis	46.15%	473	WP_107830406.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.80%	457	WP_022612328.1
nicotinate phosphoribosyltransferase	Pedobacter sp. SYP-B3415	45.76%	473	WP_129715466.1
nicotinate phosphoribosyltransferase	Variovorax sp. EL159	45.93%	481	WP_093243170.1
nicotinate phosphoribosyltransferase	Variovorax sp. RKNM96	45.83%	481	WP_206176054.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Variovorax	45.51%	481	WP_108142461.1
nicotinate phosphoribosyltransferase	Chryseolinea serpens	46.09%	471	WP_084138515.1
nicotinate phosphoribosyltransferase	Variovorax sp. OV084	45.83%	481	WP_093080925.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.80%	457	WP_022607381.1
nicotinate phosphoribosyltransferase	Variovorax sp. PAMC26660	45.51%	481	WP_187112692.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	48.84%	458	WP_196764424.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. CF118	46.17%	471	WP_090101316.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	48.84%	457	WP_023397034.1
nicotinate phosphoribosyltransferase	Synechocystis sp. PCC 6714	47.08%	460	WP_028948456.1
nicotinate phosphoribosyltransferase	Alcanivorax nanhaiticus	45.83%	470	WP_035233575.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.80%	457	WP_022598983.1

nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.59%	457	WP_022609468.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.80%	457	WP_004404065.1
nicotinate phosphoribosyltransferase	Caulobacter henricii	45.89%	465	WP_035078106.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.78%	458	WP_196766448.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.78%	458	WP_196764542.1
nicotinate phosphoribosyltransferase	Caulobacter henricii	45.89%	465	WP_035091730.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	48.41%	458	WP_211009924.1
nicotinate phosphoribosyltransferase	endosymbiont of Acanthamoeba sp. UWC8	48.63%	461	WP_038538065.1
nicotinate phosphoribosyltransferase	Halomonas sp. Y2R2	45.78%	459	WP_149286514.1
nicotinate phosphoribosyltransferase	Chitinophaga sp. MD30	46.30%	472	WP_095836864.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	47.78%	457	WP_046354660.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.59%	457	WP_045963174.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Variovorax	46.22%	481	WP_116910216.1
unnamed protein product		45.23%	472	WP_213486759.1
nicotinate phosphoribosyltransferase	Vibrio nigripulchritudo	45.80%	457	WP_022592346.1
nicotinate phosphoribosyltransferase	Synechococcus sp. PCC 7336	46.85%	459	WP_026101129.1
nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	45.64%	474	WP_078998275.1

nicotinate phosphoribosyltransferase	Microbulbifer variabilis	45.59%	460	WP_020413790.1
nicotinate phosphoribosyltransferase	Microbulbifer aggregans	45.19%	460	WP_069946600.1
nicotinate phosphoribosyltransferase	Moraxella macacae	46.22%	467	WP_009501441.1
nicotinate phosphoribosyltransferase	Vibrio sp. vnigr-6D03	46.01%	457	WP_101111920.1
nicotinate phosphoribosyltransferase	Variovorax sp. PAMC 28711	47.55%	479	WP_068630143.1
nicotinate phosphoribosyltransferase	Chryseolinea soli	45.66%	477	WP_119753759.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. dk17	44.47%	473	WP_146381491.1
nicotinate phosphoribosyltransferase	Paraflavitalea devenefica	45.23%	480	WP_167286825.1
nicotinate phosphoribosyltransferase	Leeia aquatica	44.63%	457	WP_168878251.1
nicotinate phosphoribosyltransferase	Vibrio penaeicida	45.80%	457	WP_126609696.1
nicotinate phosphoribosyltransferase	Variovorax sp. CF313	45.30%	481	WP_007838077.1
nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	45.44%	474	WP_057946734.1
nicotinate phosphoribosyltransferase	Chryseolinea flava	45.25%	475	WP_112746155.1
nicotinamide phosphoribosyltransferase-like	Dermacentor silvarum	40.80%	570	XP_037579235.1
nicotinate phosphoribosyltransferase	Acidovorax cavernicola	45.80%	481	WP_119553604.1
nicotinate phosphoribosyltransferase	Burkholderia lata	45.51%	477	WP_174936962.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. OK283	44.74%	474	WP_090649434.1

nicotinate phosphoribosyltransferase	Chitinophaga agrisoli	45.38%	473	WP_149836263.1
nicotinate phosphoribosyltransferase	Poalibacter uvarum	44.84%	460	WP_184334749.1
nicotinate phosphoribosyltransferase	Pedobacter sp. SYSU D00823	46.35%	482	WP_207534434.1
nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	45.44%	474	WP_074863778.1
nicotinate phosphoribosyltransferase	Synechocystis salina	47.26%	456	WP_194019301.1
nicotinate phosphoribosyltransferase	Microbulbifer thermotolerans	45.32%	465	WP_067150102.1
nicotinate phosphoribosyltransferase	Burkholderia sp. Ac-20392	45.51%	477	WP_206123903.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. SP1R1	45.17%	472	WP_183568439.1
nicotinate phosphoribosyltransferase	Methylomonas rhizoryzae	45.91%	462	WP_150049226.1
nicotinate phosphoribosyltransferase	Vibrio sp. S9_S30	45.38%	457	WP_190335878.1
nicotinate phosphoribosyltransferase	Aestuariuspira insulae	46.36%	475	WP_115937160.1
nicotinate phosphoribosyltransferase	Niastella sp. SYSU D00799	46.47%	491	WP_207514658.1
nicotinate phosphoribosyltransferase	Synechocystis sp. FACHB-383	47.26%	456	WP_190599906.1
nicotinate phosphoribosyltransferase	Deminuibacter soli	46.35%	484	WP_116846561.1
nicotinate phosphoribosyltransferase	Burkholderia lata	45.51%	477	WP_174985541.1
nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	45.44%	474	WP_096379383.1
nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	45.23%	474	WP_206410754.1

nicotinate phosphoribosyltransferase	Variovorax paradoxus	45.55%	481	WP_057593555.1
nicotinate phosphoribosyltransferase	Mucilaginibacter oryzae	45.13%	471	WP_109607796.1
nicotinate phosphoribosyltransferase	Lysobacter sp. Root983	45.23%	474	WP_057160250.1
nicotinate phosphoribosyltransferase	Variovorax paradoxus	45.55%	481	WP_208507474.1
nicotinate phosphoribosyltransferase	Luteimonas sp. SJ-92	44.10%	474	WP_180677306.1
nicotinate phosphoribosyltransferase	Deinococcus cellulosilyticus	46.84%	457	WP_146884252.1
nicotinate phosphoribosyltransferase	Burkholderia lata	45.30%	477	WP_174945660.1
nicotinate phosphoribosyltransferase	Variovorax sp. 1615	45.55%	481	WP_209540548.1
nicotinate phosphoribosyltransferase	Microbulbifer thermotolerans	45.11%	465	WP_074902077.1
nicotinate phosphoribosyltransferase	Luteibacter sp. 329MFSHa	43.92%	473	WP_090059091.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	46.72%	458	WP_196760576.1
nicotinate phosphoribosyltransferase	Burkholderia lata	45.09%	477	WP_174932841.1
nicotinate phosphoribosyltransferase	Lysobacter sp. Root916	45.23%	474	WP_057163575.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	45.09%	477	WP_174943287.1
nicotinate phosphoribosyltransferase	Burkholderia lata	45.09%	477	WP_175030891.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Aphanizomenonaceae	46.12%	463	WP_040553859.1
nicotinate phosphoribosyltransferase	Variovorax paradoxus	45.55%	481	WP_203203860.1

MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Burkholderia	45.09%	477	WP_166895475.1
nicotinate phosphoribosyltransferase	Mucilaginibacter mali	44.47%	473	WP_173414554.1
nicotinate phosphoribosyltransferase	Filimonas sp. YR581	45.72%	481	WP_116734937.1
nicotinate phosphoribosyltransferase	Lysobacter sp. K5869	44.40%	474	WP_215837198.1
nicotinate phosphoribosyltransferase	Lysobacter sp. cf310	45.23%	474	WP_091633503.1
nicotinate phosphoribosyltransferase	Mucilaginibacter pankratovii	44.51%	471	WP_191189719.1
nicotinate phosphoribosyltransferase	Burkholderia sp. Ac-20384	45.09%	477	WP_206144051.1
nicotinate phosphoribosyltransferase	Acaryochloris sp. RCC1774	43.79%	458	WP_110985241.1
nicotinate phosphoribosyltransferase	Lysobacter alkalisoli	44.74%	487	WP_141624649.1
nicotinate phosphoribosyltransferase	Burkholderia sp. BCC0322	44.89%	477	WP_175968842.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Burkholderia	45.09%	477	WP_069252156.1
nicotinate phosphoribosyltransferase	Variovorax sp. PDC80	45.34%	481	WP_093355202.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Caulobacter	44.12%	463	WP_110118195.1
nicotinate phosphoribosyltransferase	Mucilaginibacter psychrotolerans	44.40%	471	WP_133231431.1
nicotinate phosphoribosyltransferase	Burkholderia lata	45.09%	477	WP_174952572.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.89%	477	WP_046548351.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. OK098	44.03%	472	WP_073404934.1

nicotinate phosphoribosyltransferase	Burkholderia sp. BCC0405	44.89%	477	WP_175953292.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.68%	477	WP_124578241.1
nicotinate phosphoribosyltransferase	Lysobacter capsici	44.40%	474	WP_036110648.1
nicotinate phosphoribosyltransferase	Burkholderia sp. Se-20378	45.30%	477	WP_206134239.1
nicotinate phosphoribosyltransferase	Lysobacter capsici	44.81%	474	WP_215495265.1
nicotinate phosphoribosyltransferase	Pseudoflavitalea sp. SYSU D01084	45.60%	479	WP_205511655.1
nicotinate phosphoribosyltransferase	Lysobacter sp. Root690	44.61%	474	WP_056110948.1
nicotinate phosphoribosyltransferase	Burkholderia lata	44.89%	477	WP_175016235.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.89%	477	WP_175041117.1
nicotinate phosphoribosyltransferase	Burkholderia sp. AU6039	45.09%	477	WP_089475280.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Burkholderia	44.89%	477	WP_122476424.1
nicotinate phosphoribosyltransferase	Cellvibrio sp. YR554	45.13%	464	WP_208110304.1
nicotinate phosphoribosyltransferase	Burkholderia sp. Tr-20390	44.89%	477	WP_206149531.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. FT3.2	43.92%	472	WP_184549465.1
nicotinate phosphoribosyltransferase	Lysobacter sp. yr284	44.61%	474	WP_091794923.1
nicotinate phosphoribosyltransferase	Foetidibacter luteolus	45.88%	480	WP_153798260.1
nicotinate phosphoribosyltransferase	Burkholderia lata	44.89%	477	WP_175044317.1

nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	44.61%	474	WP_207523671.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. SG538B	45.55%	472	WP_176627709.1
nicotinate phosphoribosyltransferase	Burkholderia sp. AU31652	44.89%	477	WP_089433005.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.89%	477	WP_124494750.1
nicotinate phosphoribosyltransferase	Mucilaginibacter gotjawali	44.24%	471	WP_096356197.1
nicotinate phosphoribosyltransferase	Vibrio penaeicida	45.17%	457	WP_104400876.1
nicotinate phosphoribosyltransferase	Chitinivorax sp. B	44.94%	459	WP_137937993.1
nicotinate phosphoribosyltransferase	Synechocystis sp. LEGE 06083	46.84%	456	WP_194072815.1
nicotinate phosphoribosyltransferase	Burkholderia lata	44.89%	477	WP_011354434.1
nicotinate phosphoribosyltransferase	Lysobacter enzymogenes	45.34%	474	WP_123649136.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.89%	477	WP_124619061.1
nicotinate phosphoribosyltransferase	Variovorax sp. dw_954	45.84%	482	WP_213952862.1
nicotinate phosphoribosyltransferase	Burkholderia sp. Ac-20344	45.09%	477	WP_205786422.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.68%	477	WP_175842282.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Burkholderia	44.89%	477	WP_089441950.1
nicotinate phosphoribosyltransferase	Burkholderia sp. HI2714	44.89%	477	WP_089485327.1
nicotinate phosphoribosyltransferase	Lysobacter sp. Root604	44.81%	474	WP_056178160.1

MULTISPECIES: nicotinate phosphoribosyltransferase	Burkholderia	44.89%	477	WP_089458617.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Mucilaginibacter	45.55%	472	WP_129571949.1
nicotinate phosphoribosyltransferase	Burkholderia contaminans	44.68%	477	WP_175812769.1
nicotinate phosphoribosyltransferase	Ralstonia sp. ASV6	45.72%	477	WP_199031363.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Burkholderia cepacia complex	44.89%	477	WP_175239396.1
nicotinate phosphoribosyltransferase	Lysobacter antibioticus	44.81%	474	WP_057918818.1
nicotinate phosphoribosyltransferase	Luteibacter rhizovincinus	45.02%	472	WP_132145436.1
nicotinate phosphoribosyltransferase	Luteibacter sp. UNC138MFC05.1	43.60%	473	WP_090175555.1
nicotinate phosphoribosyltransferase	Xanthomonas sp. GPE 39	44.81%	474	WP_045728230.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. NFR10	44.51%	472	WP_090526283.1
nicotinate phosphoribosyltransferase	Mucilaginibacter sp. YR332	44.31%	472	WP_121230029.1
nicotinate phosphoribosyltransferase	Pseudoalteromonas luteoviolacea	46.51%	458	WP_211036465.1
nicotinate phosphoribosyltransferase	Gynuella sunshinyii	45.86%	458	WP_044616760.1
nicotinate phosphoribosyltransferase	Burkholderia sp. Se-20373	44.47%	477	WP_206139413.1
nicotinate phosphoribosyltransferase	Ralstonia solanacearum	44.47%	483	WP_119447796.1
nicotinate phosphoribosyltransferase	Moraxella osloensis	44.75%	468	WP_144918487.1
MULTISPECIES: nicotinate phosphoribosyltransferase	unclassified Lysobacter	44.61%	474	WP_215056362.1

nicotinate phosphoribosyltransferase	Lysobacter antibioticus	44.61%	474	WP_031373421.1
MULTISPECIES: nicotinate phosphoribosyltransferase	Burkholderia	44.89%	477	WP_089462279.1