

Table S9D. Transporters in the NP-SAG (*Ca. Nitromaritima* sp. NP) genome.

| Function TC | ORF Name | Family ID | Transporter Family | Subfamily | Substrate/Function |
|-------------|--------------|------------|---|-----------------|---|
| 3.A.1 | Nspl16_00034 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | phosphonate |
| 3.A.1 | Nspl16_00035 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipoprotein |
| 3.A.1 | Nspl16_00036 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | cell division |
| 2.A.64 | Nspl16_00059 | Tat | The Twin Arginine Targeting (Tat) Family | | protein export |
| 3.A.1 | Nspl16_00085 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipoprotein |
| 3.A.1 | Nspl16_00086 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | efflux (antimicrobial peptide?) |
| 2.A.66 | Nspl16_00124 | MOP | The Multidrug/Oligosaccharidyl-lipid/Poly saccharide (MOP) Flippase Superfamily | MVF | virulence factor MviN |
| 9.A.19 | Nspl16_00136 | MgtE | The Mg ²⁺ Transporter-E (MgtE) Family | | magnesium ion |
| 3.A.1 | Nspl16_00167 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipid A |
| 2.A.7 | Nspl16_00171 | DMT | The Drug/Metabolite Transporter (DMT) Superfamily | | drug/metabolite? |
| 3.A.1 | Nspl16_00193 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | daunorubicin |
| 4.A | Nspl16_00196 | SSPTS | Sugar Specific PTS | EnzymeIA | fructose |
| 4.A | Nspl16_00198 | GPTS | General PTS | EnzymeI | |
| 3.A.1 | Nspl16_00205 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding protein | oligopeptide |
| 3.A.1 | Nspl16_00206 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | dipeptide/oligopeptide |
| 3.A.1 | Nspl16_00207 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | dipeptide/oligopeptide |
| 3.A.1 | Nspl16_00208 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | glycine betaine |
| 2.A.21 | Nspl16_00234 | SSS | The Solute:Sodium Symporter (SSS) Family | | sodium ion:proline symporter |
| 2.A.66 | Nspl16_00238 | MOP | The Multidrug/Oligosaccharidyl-lipid/Poly saccharide (MOP) Flippase Superfamily | PST | polysaccharide export |
| 1.A.46 | Nspl16_00242 | Bestrophin | The Anion Channel-forming Bestrophin (Bestrophin) Family | | Bestrophin anion channel |
| 3.A.15 | Nspl16_00265 | MTB | The Outer Membrane Protein Secretory Main Terminal Branch (MTB) | | |
| 2.A.1 | Nspl16_00399 | MFS | The Major Facilitator Superfamily (MFS) | | nitrate |
| 3.A.1 | Nspl16_00412 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | glycine betaine |
| 2.A.9 | Nspl16_00455 | Oxa1 | The Cytochrome Oxidase Biogenesis (Oxa1) Family | | 60 KD inner membrane protein OxaA homolog |
| 2.A.47 | Nspl16_00469 | DASS | The Divalent Anion:Na ⁺ Symporter (DASS) Family | | sodium ion:dicarboxylate/sulfate |
| 2.A.1 | Nspl16_00486 | MFS | The Major Facilitator Superfamily (MFS) | | multidrug efflux |
| 3.A.1 | Nspl16_00490 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | dipeptide/oligopeptide |
| 3.A.1 | Nspl16_00491 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | dipeptide/oligopeptide |
| 3.A.1 | Nspl16_00492 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | phosphonate |
| 9.A.6 | Nspl16_00499 | ATP-E | The ATP Exporter (ATP-E) Family | | |
| 2.A.64 | Nspl16_00521 | Tat | The Twin Arginine Targeting (Tat) Family | | protein export |
| 3.A.10 | Nspl16_00533 | H+-PPase | The H ⁺ -translocating Pyrophosphatase (H ⁺ -PPase) Family | | proton (vacuolar) |
| 3.A.1 | Nspl16_00556 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | toluene tolerance |
| 2.A.22 | Nspl16_00560 | NSS | The Neurotransmitter:Sodium Symporter (NSS) Family | | sodium ion:solute symporter |
| 3.A.1 | Nspl16_00571 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipoprotein |
| 3.A.1 | Nspl16_00572 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | lipoprotein releasing |
| 3.A.1 | Nspl16_00573 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | lipoprotein releasing |
| 3.A.1 | Nspl16_00579 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | iron-hydroxamate |
| 3.A.1 | Nspl16_00647 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | dipeptide/oligopeptide |
| 3.A.1 | Nspl16_00648 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | dipeptide/oligopeptide |
| 3.A.1 | Nspl16_00672 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | leucine/valine |
| 3.A.6 | Nspl16_00680 | IIISP | The Type III (Virulence-related) Secretory Pathway (IIISP) Family | | |
| 2.A.50 | Nspl16_00692 | GUP | The Glycerol Uptake (GUP) Family | | glycerol:proton symporter |
| 2.A.1 | Nspl16_00733 | MFS | The Major Facilitator Superfamily (MFS) | | multidrug efflux? |
| 2.A.28 | Nspl16_00743 | BASS | The Bile Acid:Na ⁺ Symporter (BASS) Family | | sodium ion/? |
| 2.A.19 | Nspl16_00752 | CaCA | The Ca ²⁺ :Cation Antiporter (CaCA) Family | | sodium ion:calcium ion antiporter |
| 2.A.41 | Nspl16_00759 | CNT | The Concentrative Nucleoside Transporter (CNT) Family | | sodium ion:nucleoside symporter |
| 1.A.30 | Nspl16_00764 | Mot/Exb | The H ⁺ - or Na ⁺ -translocating Bacterial Flagellar Motor 1ExbBD Outer Membrane Transport Energizer (Mo) | | |
| 1.B.6 | Nspl16_00766 | OOP | The OmpA-OmpF Porin (OOP) Family | | |
| 2.A.1 | Nspl16_00767 | MFS | The Major Facilitator Superfamily (MFS) | | Acetyl-CoA:CoA antiporter |
| 2.A.64 | Nspl16_00777 | Tat | The Twin Arginine Targeting (Tat) Family | | protein export |
| 3.A.1 | Nspl16_00812 | ABC | The ATP-binding Cassette (ABC) Superfamily | ABC+ membrane | multidrug |
| 1.B.6 | Nspl16_00822 | OOP | The OmpA-OmpF Porin (OOP) Family | | |
| 1.A.30 | Nspl16_00823 | Mot/Exb | The H ⁺ - or Na ⁺ -translocating Bacterial Flagellar Motor 1ExbBD Outer Membrane Transport Energizer (Mo) | | |
| 3.A.1 | Nspl16_00824 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | glycine betaine |
| 3.A.1 | Nspl16_00825 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | phosphate |
| 2.A.4 | Nspl16_00827 | CDF | The Cation Diffusion Facilitator (CDF) Family | | cation efflux |
| 9.A.10 | Nspl16_00836 | ILT | The Iron/Lead Transporter (ILT) Superfamily | | iron ion |
| 2.A.25 | Nspl16_00860 | AGCS | The Alanine or Glycine:Cation Symporter (AGCS) Family | | sodium ion:alanine symporter |
| 1.A.30 | Nspl16_00874 | Mot/Exb | The H ⁺ - or Na ⁺ -translocating Bacterial Flagellar Motor 1ExbBD Outer Membrane Transport Energizer (Mo) | | |
| 3.A.1 | Nspl16_00878 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | cobalt |
| 1.A.1 | Nspl16_00887 | VIC | The Voltage-gated Ion Channel (VIC) Superfamily | | potassium ion channel |
| 2.A.6 | Nspl16_00894 | RND | The Resistance-Nodulation-Cell Division (RND) Superfamily | HAE3 | multidrug/solvent efflux (HAE3 subfamily) |
| 3.A.1 | Nspl16_01010 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | efflux (antimicrobial peptide?) |
| 3.A.1 | Nspl16_01011 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | efflux (antimicrobial peptide?) |
| 3.A.1 | Nspl16_01012 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipoprotein |
| 2.A.6 | Nspl16_01013 | RND | The Resistance-Nodulation-Cell Division (RND) Superfamily | | multidrug efflux |
| 9.A.4 | Nspl16_01028 | YggT | The YggT or Fancf1 K ⁺ Uptake-B (FkbB; YggT) Family | | ? |
| 3.A.1 | Nspl16_01045 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | ? (Fe-S assembly/SufBCD system) |
| 3.A.1 | Nspl16_01046 | ABC | The ATP-binding Cassette (ABC) Superfamily | ABC | ? (Fe-S assembly/SufBCD system) |
| 3.A.1 | Nspl16_01047 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | ? (Fe-S assembly/SufBCD system) |
| 3.A.1 | Nspl16_01057 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | phosphate |
| 3.A.1 | Nspl16_01058 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | phosphonate |
| 3.A.1 | Nspl16_01059 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | branched-chain amino acid |
| 3.A.1 | Nspl16_01060 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | branched-chain amino acid |
| 3.A.1 | Nspl16_01061 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | branched-chain amino acid |
| 3.A.1 | Nspl16_01062 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | urea |
| 2.A.20 | Nspl16_01078 | PIT | The Inorganic Phosphate Transporter (PIT) Family | | phosphate |
| 2.A.6 | Nspl16_01082 | RND | The Resistance-Nodulation-Cell Division (RND) Superfamily | HME | cobalt/zinc/cadmium ion efflux (HME subfamily) |
| 3.A.7 | Nspl16_01099 | IVSP | The Type IV (Conjugal DNA-Protein Transfer or VirB) Secretory Pathway (IVSP) Family | | |
| 1.A.11 | Nspl16_01109 | Amt | The Ammonia Transporter Channel (Amt) Family | | ammonium |
| 2.A.64 | Nspl16_01115 | Tat | The Twin Arginine Targeting (Tat) Family | | protein export |
| 2.A.1 | Nspl16_01185 | MFS | The Major Facilitator Superfamily (MFS) | | multidrug efflux? |
| 2.A.40 | Nspl16_01200 | NCS2 | The Nucleobase:Cation Symporter-2 (NCS2) Family | | xanthine/uracil |
| 2.A.6 | Nspl16_01217 | RND | The Resistance-Nodulation-Cell Division (RND) Superfamily | SecDF | protein export (SecDF) |
| 3.A.1 | Nspl16_01223 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | lipoprotein releasing |
| 2.A.21 | Nspl16_01250 | SSS | The Solute:Sodium Symporter (SSS) Family | | sodium ion:proline symporter |
| 3.A.1 | Nspl16_01276 | ABC | The ATP-binding Cassette (ABC) Superfamily | ABC+ membrane | multidrug |
| 2.A.23 | Nspl16_01292 | DAACS | The Dicarboxylate/Amino Acid:Cation (Na ⁺ or H ⁺) Symporter (DAACS) Family | | proton/sodium ion:glutamate/aspartate symporter |
| 2.A.1 | Nspl16_01301 | MFS | The Major Facilitator Superfamily (MFS) | | multidrug efflux |
| 3.A.1 | Nspl16_01303 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding protein | ? |
| 3.A.15 | Nspl16_01316 | MTB | The Outer Membrane Protein Secretory Main Terminal Branch (MTB) | | |
| 3.A.15 | Nspl16_01317 | MTB | The Outer Membrane Protein Secretory Main Terminal Branch (MTB) | | |
| 3.A.15 | Nspl16_01318 | MTB | The Outer Membrane Protein Secretory Main Terminal Branch (MTB) | | |

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| 3.A.1 | NspL16_01333 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | lipoprotein releasing |
| 3.A.1 | NspL16_01334 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipoprotein |
| 3.A.1 | NspL16_01338 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipid A |
| 3.A.15 | NspL16_01342 | MTB | The Outer Membrane Protein Secreting Main Terminal Branch (MTB) | | |
| 2.A.49 | NspL16_01401 | CIC | The Chloride Carrier/Channel (CIC) Family | | chloride ion channel |
| 3.A.3 | NspL16_01404 | P-ATPase | The P-type ATPase (P-ATPase) Superfamily | | zinc/cadmium/cobalt ion |
| 2.A.1 | NspL16_01437 | MFS | The Major Facilitator Superfamily (MFS) | | multidrug efflux |
| 3.A.6 | NspL16_01446 | IIISP | The Type III (Virulence-related) Secretory Pathway (IIISP) Family | | |
| 3.A.6 | NspL16_01447 | IIISP | The Type III (Virulence-related) Secretory Pathway (IIISP) Family | | |
| 3.A.6 | NspL16_01448 | IIISP | The Type III (Virulence-related) Secretory Pathway (IIISP) Family | | |
| 3.A.6 | NspL16_01449 | IIISP | The Type III (Virulence-related) Secretory Pathway (IIISP) Family | | |
| 3.A.6 | NspL16_01450 | IIISP | The Type III (Virulence-related) Secretory Pathway (IIISP) Family | | |
| 2.A.6 | NspL16_01479 | RND | The Resistance-Nodulation-Cell Division (RND) Superfamily | HAE3 | multidrug/solvent efflux (HAE3 subfamily) |
| 2.A.64 | NspL16_01481 | Tat | The Twin Arginine Targeting (Tat) Family | | protein export |
| 3.A.1 | NspL16_01514 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | phosphonate |
| 9.A.40 | NspL16_01550 | HCC | The HlyC/CorC (HCC) Family | | hemolysin C (HlyC) homolog |
| 9.A.40 | NspL16_01551 | HCC | The HlyC/CorC (HCC) Family | | hemolysin C (HlyC) homolog |
| 3.A.1 | NspL16_01576 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | iron-hydroxamate |
| 3.A.1 | NspL16_01577 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | phosphonate |
| 3.A.1 | NspL16_01578 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | cobalamin/Fe ³⁺ -siderophores |
| 2.A.37 | NspL16_01593 | CPA2 | The Monovalent Cation:Proton Antiporter-2 (CPA2) Family | | potassium/sodium ion:proton antiporter |
| 3.A.1 | NspL16_01599 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | lipoprotein |
| 2.A.43 | NspL16_01607 | LCT | The Lysosomal Cystine Transporter (LCT) Family | | |
| 2.A.6 | NspL16_01676 | RND | The Resistance-Nodulation-Cell Division (RND) Superfamily | HAE3 | multidrug/solvent efflux (HAE3 subfamily) |
| 3.A.1 | NspL16_01701 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding protein | oligopeptide |
| 2.A.50 | NspL16_01739 | GUP | The Glycerol Uptake (GUP) Family | | glycerol:proton symporter |
| 9.A.19 | NspL16_01851 | MgtE | The Mg ²⁺ Transporter-E (MgtE) Family | | magnesium ion |
| 1.A.23 | NspL16_01854 | MscS | The Small Conductance Mechanosensitive Ion Channel (MscS) Family | | small-conductance mechanosensitive ion channel |
| 1.A.23 | NspL16_01882 | MscS | The Small Conductance Mechanosensitive Ion Channel (MscS) Family | | small-conductance mechanosensitive ion channel |
| 1.A.1 | NspL16_01931 | VIC | The Voltage-gated Ion Channel (VIC) Superfamily | | potassium ion channel |
| 3.A.1 | NspL16_01946 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | ? |
| 1.A.34 | NspL16_02023 | EVE1-C | The Envelope Virus E1 Channel (EVE1-C) Family | | |
| 9.A.19 | NspL16_02034 | MgtE | The Mg ²⁺ Transporter-E (MgtE) Family | | magnesium ion |
| 2.A.21 | NspL16_02062 | SSS | The Solute:Sodium Symporter (SSS) Family | | sodium ion:proline symporter |
| 2.A.47 | NspL16_02166 | DASS | The Divalent Anion:Na ⁺ Symporter (DASS) Family | | sodium ion:dicarboxylate/sulfate |
| 2.A.21 | NspL16_02188 | SSS | The Solute:Sodium Symporter (SSS) Family | | sodium ion:proline symporter |
| 3.A.1 | NspL16_02217 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | molybdate |
| 3.A.1 | NspL16_02218 | ABC | The ATP-binding Cassette (ABC) Superfamily | membrane | molybdate |
| 3.A.1 | NspL16_02219 | ABC | The ATP-binding Cassette (ABC) Superfamily | binding | molybdenate |
| 2.A.44 | NspL16_02239 | FNT | The Formate-Nitrite Transporter (FNT) Family | | formate/nitrite |
| 3.A.1 | NspL16_02263 | ABC | The ATP-binding Cassette (ABC) Superfamily | ABC | ? (Uup homolog/duplicated ATPase) |