

Supplementary materials

Title: Molecular evolution of gas cavity in [NiFeSe] hydrogenases resurrected *in silico*

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supplementary Figures S1 to S9

supplementary Table S1

Supplementary Fig. S1

>*D. vulgaris* Hildenborough

GACCCGTGActgggctgtgccgtgcacgtgctgcacgctgagtcggcaaggtcgcgtcatcg
aagtgaagtag

$\Delta G = -27.70$ kJ/mol

```

          10          20
          | CG A       gu c     cg
--GACC UG cugggcu gc gugca \
cugg ac ggccuga cg cacgu u
\ ^ a- -      gu -     cg
  50          40          30
          60
gccg-   c
ucau g
agug a
gauga   a
  70

```

>*D. vulgaris* Miyazaki F

GACCCGTGActgggctgtgccgtgcacgtgctgcacgctgaatccggcaaggctccgtcgtcg
aagtgaagtag

$\Delta G = -22.80$ kJ/mol

```

          10          20          30
GACC----- - u- g ug-- -| c
          CG UGAc gggcu ugccg ca cgug \
          gc gcug ucugg acggc gu gcac u
gaugaagugaa u cc a cuaa c^ g
  70       60       50       40

```

>*D. alaskensis* G20

GACCCGTGActgggctgcgcgtgcacgtgctgcacgcagagtccggtaaggttccgtagtcg
aattctag

$\Delta G = -23.90$ kJ/mol

```

          10          20
----- -----| GUG       gc c     cg
          GAC       CC  Acugggcu gc gugca \
          cug       gg  uggccuga cg cacgu u
gaucuuuaag augccuuu^ aa-      ga -     cg
  70       60       50       40       30

```

Supplementary Fig. S1

>*D. piger*

GACCCGTGActgggctgtgccgtgcacgtgctgcacgctgagaccggcgaagaatccatcgta
acctgggctag

$\Delta G = -20.80 \text{ kJ/mol}$

```

          10      20
GA-- .-GUGA| g   u   c   c
             CCC   cu ggc gugc gug a
             ggg   ga ucg cacg cgu c
gauc \ ----^ g   -   u   g
    70      40      30
          50
accg-- aga
      gcga   a
      ugcu   u
uccaag   acc
    60

```

> *B. wadsworthia* 3-1-6

GACCCGTGActgggttgtgccgtgcacgtgctgcacgccgaatccggcaaggttgcgtcatca
atatgattaa

$\Delta G = -23.50 \text{ kJ/mol}$

```

          10      20
|   CG  A       g   c   cg
--GACC  UG cuggguu ug cgugca \
  uugg  ac ggccuaa gc gcacgu u
\ ^   a- -       - c       cg
    50      40      30
          60
g---- cau
      ccgu \
      gguu   c
aauua   uaa
    70

```

>*D. baculatum*

GACCCGTGActgggctgtgccgtgcacgtgctgcacgctgagaccggtaagaacaacgttgtca
acattgactaa

$\Delta G = -21.50 \text{ kJ/mol}$

```

          10      20
----- C- -----|   g   gu   c   cg
      GAC  CGUG      Acugg cu   gc gugca \
      cug  gcac      uggcc ga   cg cacgu u
aaucaguuacaa   uu   aagaag^   a   gu   -   cg
    70      60      50      40      30

```

Supplementary Fig. S1

>*D. salexigens*

GACCCGTGActgggttgcgcgtgcacgtgctgcacgcagagaccggtaagaacatgttgttc
acgtaggcgaagggttgcataa

$\Delta G = -26.80$ kJ/mol

```

      10          20
GACCC UG .-g   g-| c     cg
      G Acu   gguu   ugc gugca \
      c ugg   ccag   acg cacgu u
aau-- gu   \ - ag^ - cg
      80          40          30
                           50
      g----- g   c
                  gugaa aa a
                  cacuu uu u
aagcggaug      g   g
      70          60

```

>*D. africanus* Walvis Bay

GATCCGTGActgggttgcgcgtgcacgtgctgagactggcgaaaagaccgttgtca
ctgtcgagtag

$\Delta G = -22.40$ kJ/mol

```

      10          20          30
----- CC   ug-   g--- - g- -| u
      GAU GUGAc   gguu   cgcc gu   ca cgug \
      cug   cacug   ccag   gcgg ca   gu gcac u
gaugag   u-   uug   aaaa   u   ga   c^   g
      70          60          50          40

```

>*D. hydrothermalis*

GATCCCATGActgggttgcgcgtgcacgtgctgcacgcagagaccggtaagagcatgtttacc
acgtggagaaggctgctag

$\Delta G = -26.40$ kJ/mol

```

      10          20
.-GAUCCAUG|   g   g   c     cg
      Acugg cu ugc gugca \
      uggcc ga acg cacgu u
\ -----^ a   g   -   cg
      40          30
      50          60
      ga -   aug   a   a
      ag   agc   uuu cc c
      uc   ucg   aga gg g
      ga   g   ga-   g   u
      80          70

```

Supplementary Fig. S1

>*D. postgatei* 2ac9

GACCCTGActgggctgcgccgtccacgtgctggacgcagataccggcaagcaaatcaaagtgg
aagtccctgtaa

$\Delta G = -17.10 \text{ kJ/mol}$

```

          10          20
.-GACCCC A g--| gcc      cg
      UG cugg   cugc    gucca \
      ac ggcc   gacg    caggu u
\ ----- - aua^ --- cg
          40          30
          50          60
      agcaaaucaa u aa
      ag gg \
      uc cc g
aaug----- c uu
          70

```

>*D. phosphitoxidans*

GACCCTGActgggttgcgccgtccacgtgctggacgtagaaaccggtcggaccgttaaagttag
acgtacccttataa

$\Delta G = -23.00 \text{ kJ/mol}$

```

          10          20
.-GACCC| guugcgc      cg
      CUGAcugg      cgucca \
      ggcuggcc      gcaggu u
\ -----^ aaagau- cg
          50          40          30
          60
      ac----- aa
      cguu a
      gcag g
aaauuuucccau au
          70

```

>*D. autrotrophicum* HRM2

GACCCCATGActgggctgtgccgtcacgtgctgcacgcagagacaggcaaaacaacgttgg
agattcccgctctag

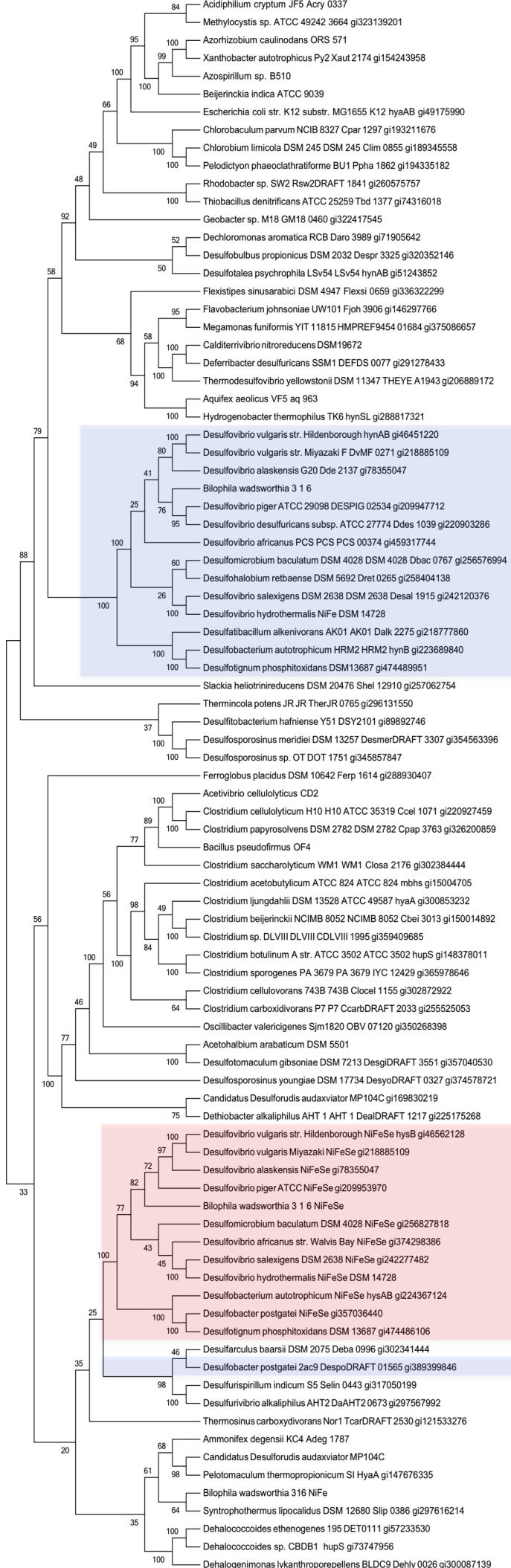
$\Delta G = -18.70 \text{ kJ/mol}$

```

          10          20
.-GACCCA| A gg g c      cg
      UG cug cu ugc gugca \
      ac gac ga acg cacgu u
\ -----^ g a- g - cg
          40          30
          50          60
      aaacaaacguuguu au
      gag u
      cuc c
gau----- gc
          70

```

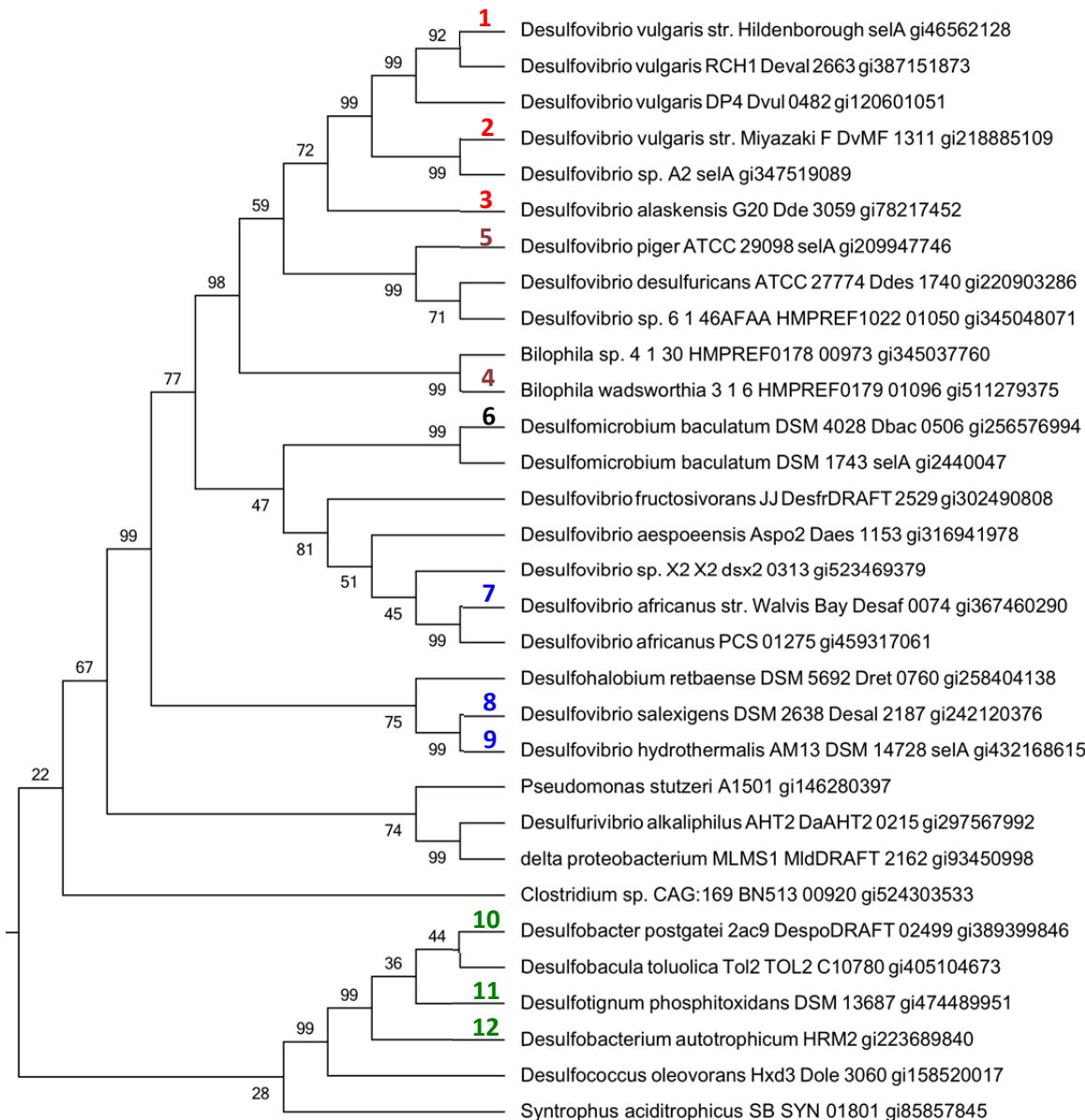
Supplementary Fig. S1: Structural models of the SECIS element predicted from the sequence immediately after the in-frame opal (UGA) codon. The stabilizing energy and secondary mRNA structure were computed on the mFold web server:
<http://mfold.rna.albany.edu/>



Supplementary Fig. S2: The phylogeny of 88 [NiFe(Se)] Hases inferred by the ML method.

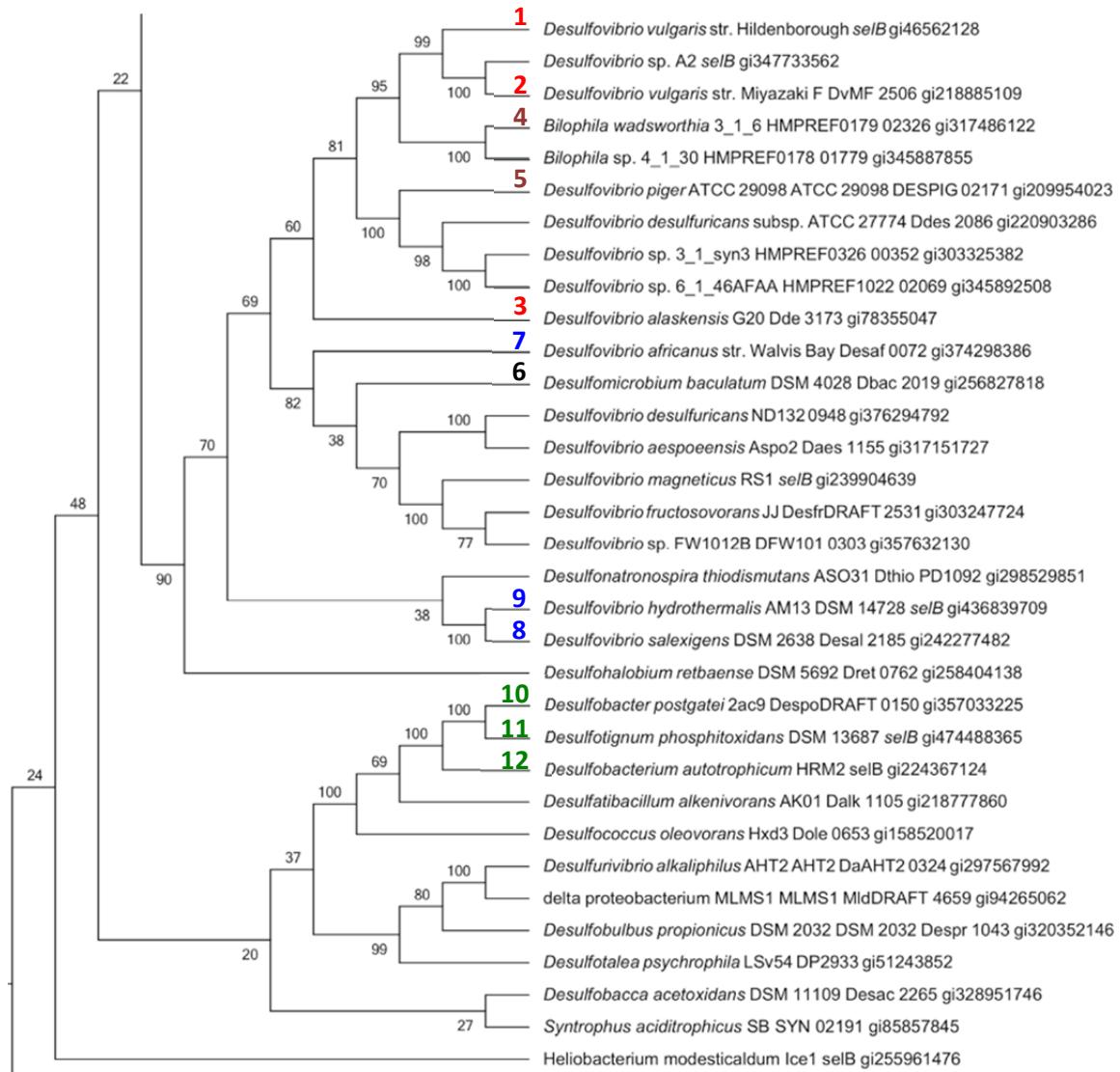
Below the branches are ML bootstrap percentages. Clades containing [NiFe] and [NiFeSe] Hases are shown in blue and red boxes, respectively.

Supplementary Fig. S3



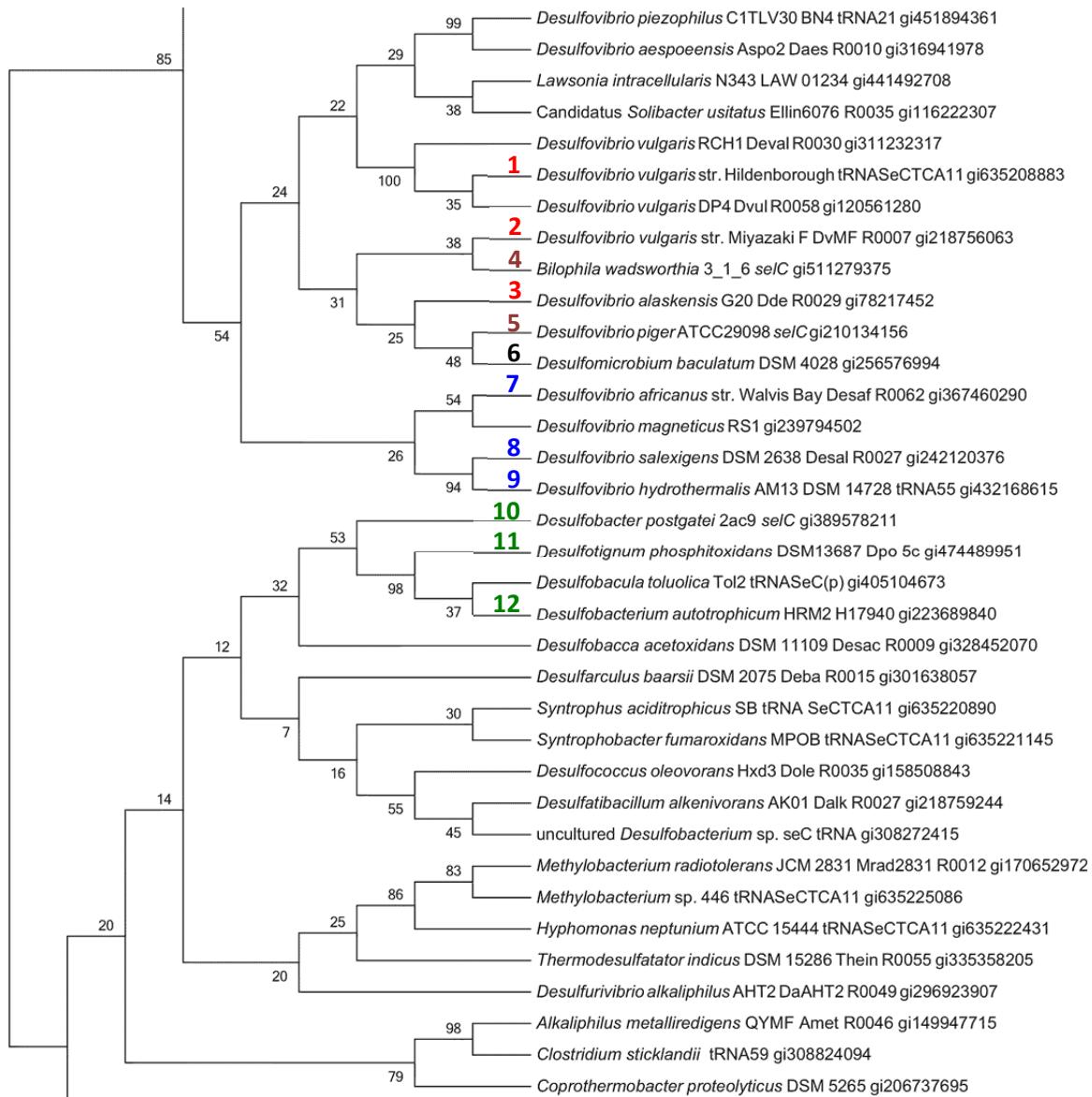
Supplementary Fig. S3: Phylogenetic relationships of selA genes in a maximum-likelihood phylogenetic tree made from 74 homologous lineages. Genes from sulphate-reducing bacteria of terrestrial origin (1-3), from clinical isolates from human faeces (4-5), of marine vent origin (7-9), and of sediment origin (10-12) are numbered in colors red, brown, blue, and green, respectively. Scores designated at each branch represent the percent of the 300-times Bootstrap test.

supplementary Fig. S4



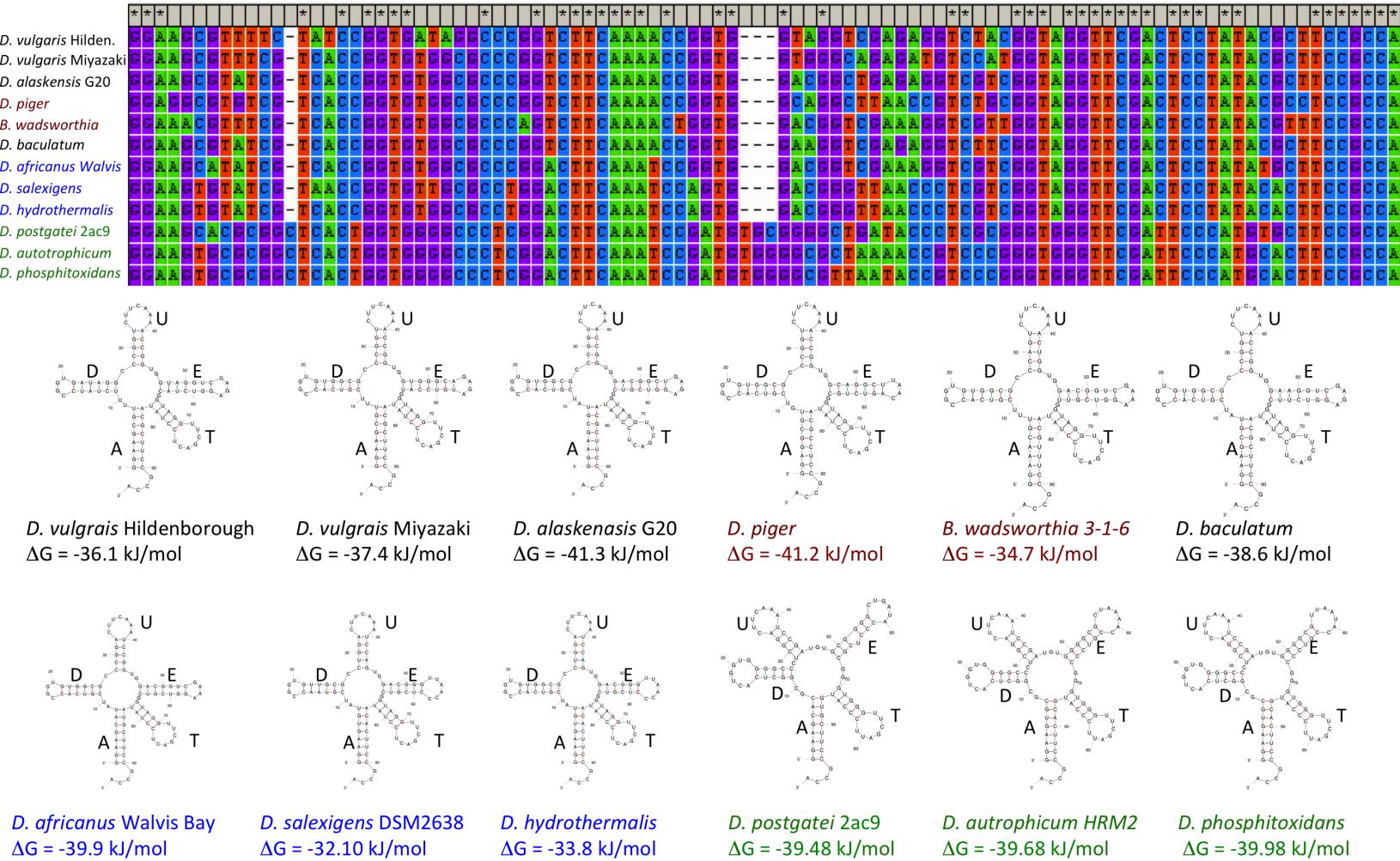
Supplementary Fig. S4: Phylogenetic relationships among the *seB* genes in a maximum-likelihood phylogenetic tree made from 88 homologous lineages. Genes from sulphate-reducing bacteria of terrestrial origin (1-3), from clinical isolates from human faeces (4-5), of marine vent origin (7-9), and of sediment origin (10-12) are numbered in colors red, brown, blue, and green, respectively. Scores designated at each branch represent the percent of the 300-times Bootstrap test.

Supplementary Fig. S5



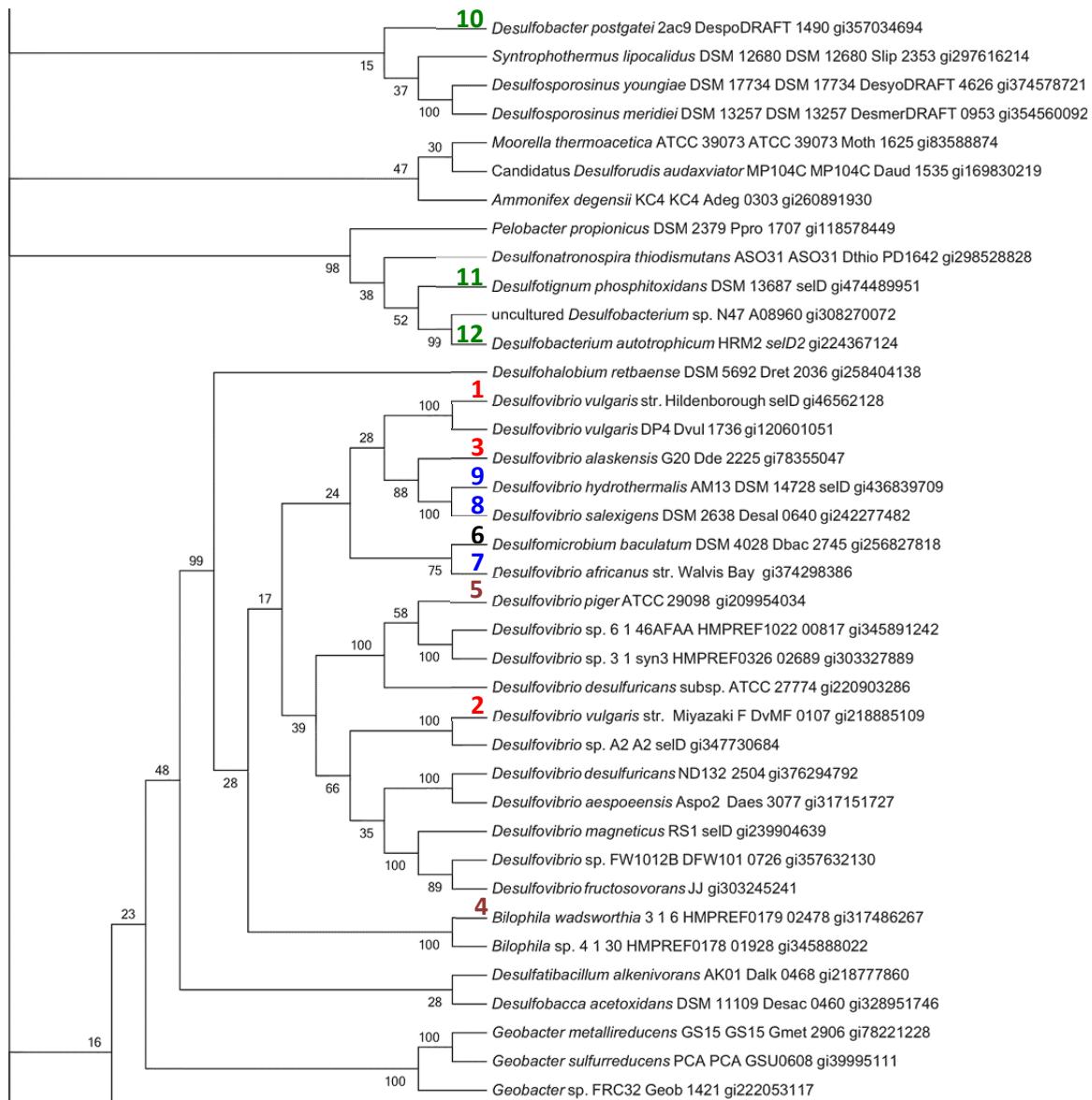
Supplementary Fig. S5: Phylogenetic relationships of *seIC* gene in the maximum-likelihood phylogenetic tree made from 48 homologous lineages. Genes from sulphate-reducing bacteria of terrestrial origin (1-3), from clinical isolates from human feaces (4-5), of marine vent origin (7-9), and of sediment origin (10-12) are numbered in colors red, brown, blue and green, respectively. Scores designated at each branch represent the percent of the 500-times Bootstrap test. The cut-off value for the consensus tree was set at a bootstrap value of 15%. Accordingly, sequences encoding *seIC* were hard to discern their phylogenetic relationships except for the three linages (10-12) due to the extra 3 bases that they have between the E and T arms of their predicted cloverleaf model.

Supplementary Fig. S6



Supplementary Fig. S6: Sequence alignment and cloverleaf models for the tRNA^{UGA} sequences encoded in genomes harboring [NiFeSe] Hase genes.

supplementary Fig. S7



Supplementary Fig. S7: Phylogenetic relationships of the *selD* genes in a maximum-likelihood phylogenetic tree made from 95 homologous lineages. Genes from sulphate-reducing bacteria of terrestrial origin (1-3), from clinical isolates from human faeces (4-5), of marine vent origin (6-9), and of sediment origin (10-12) are numbered in colors red, brown, blue, and green respectively. Scores designated at each branch represent the percent of the 300-times Bootstrap test.

Fig. S8

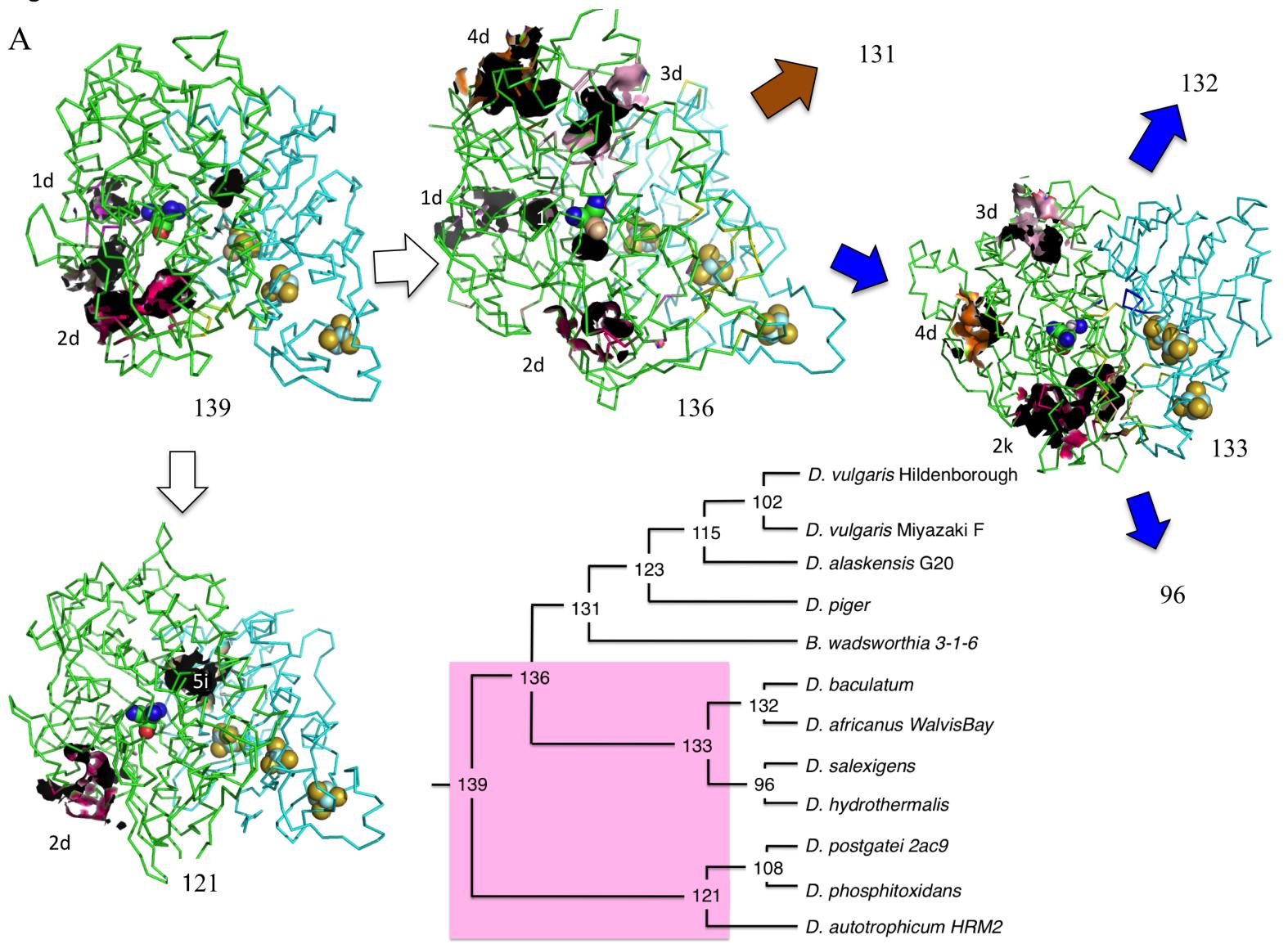


Fig. S8

B

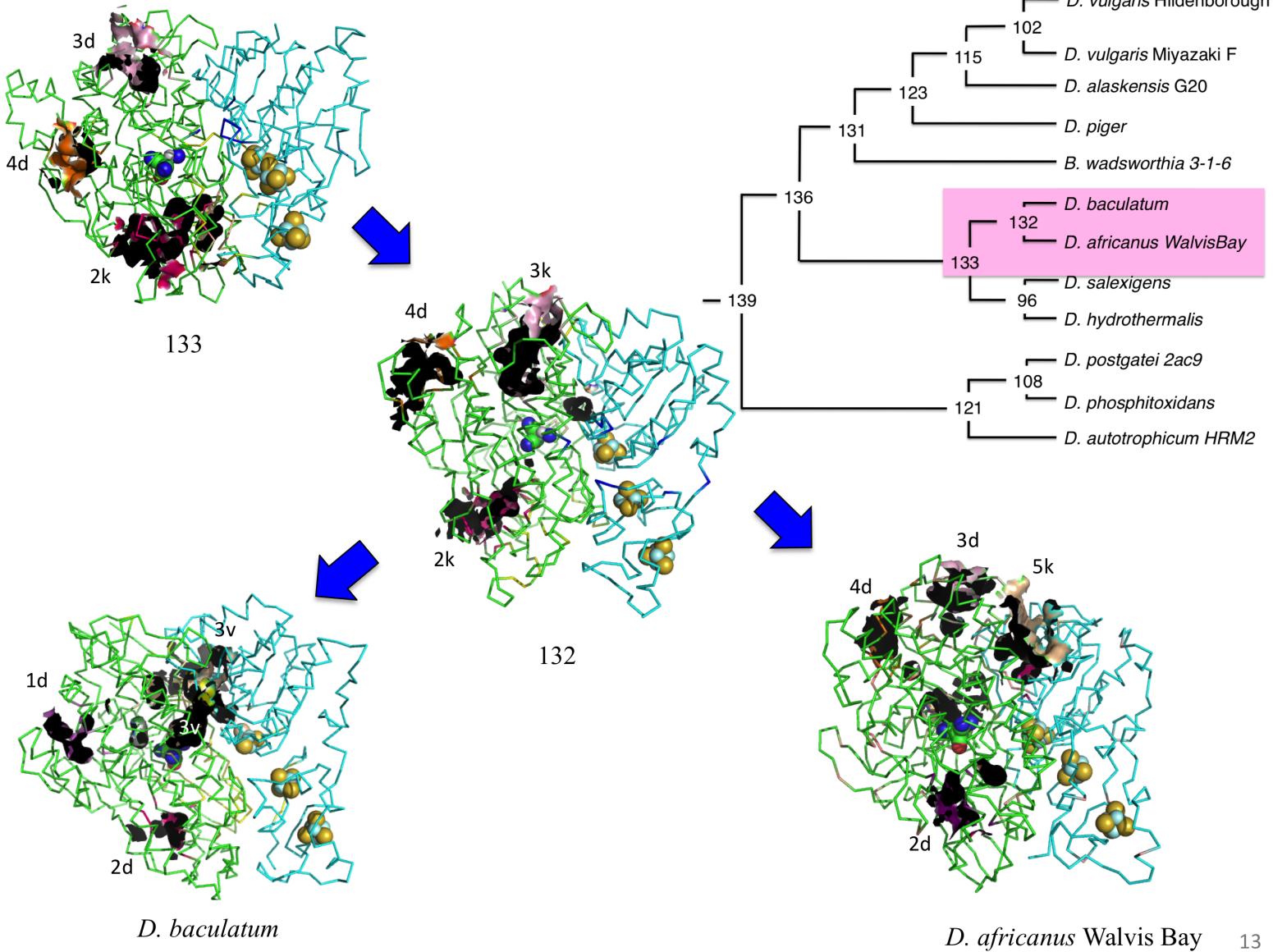


Fig. S8

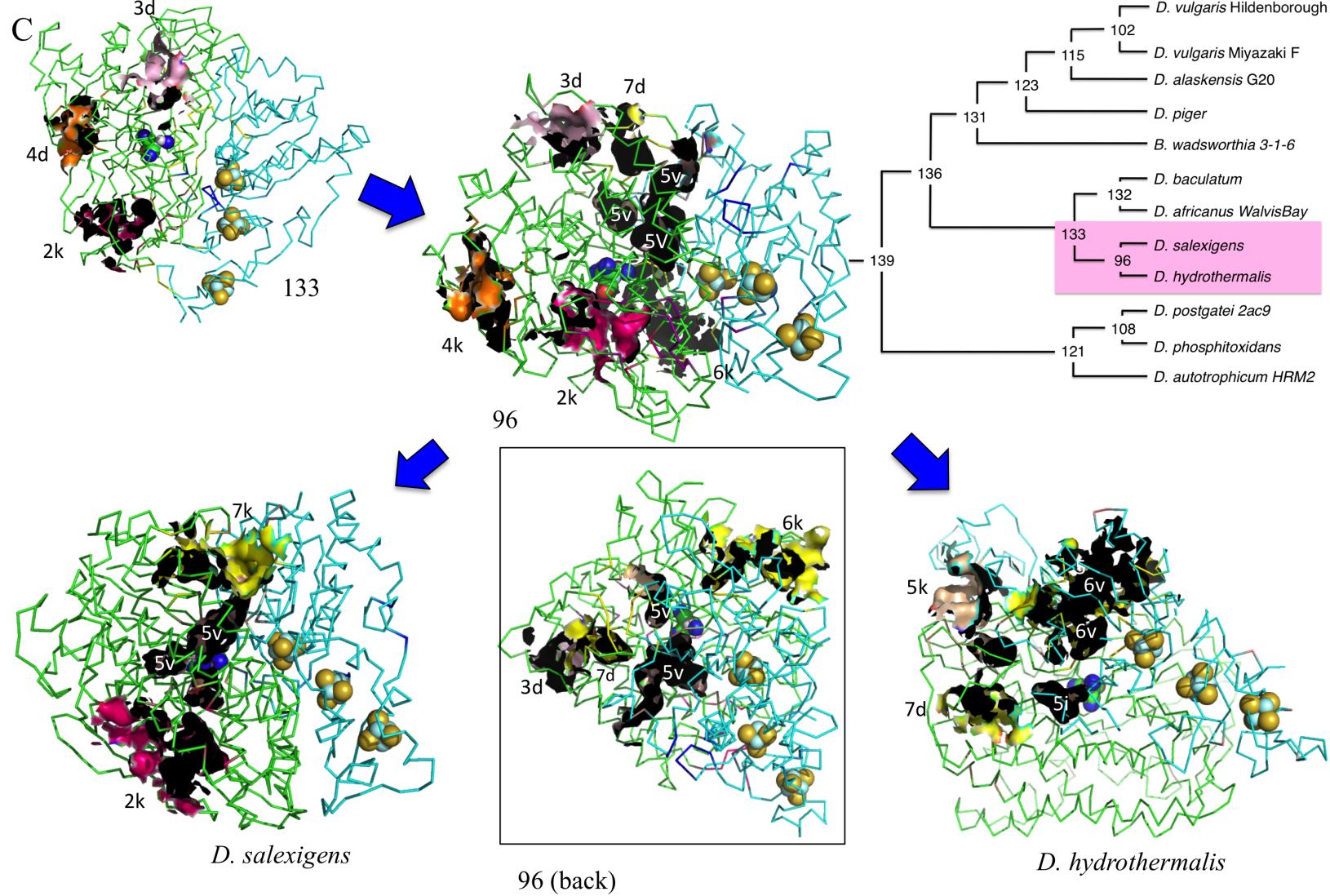


Fig. S8

D

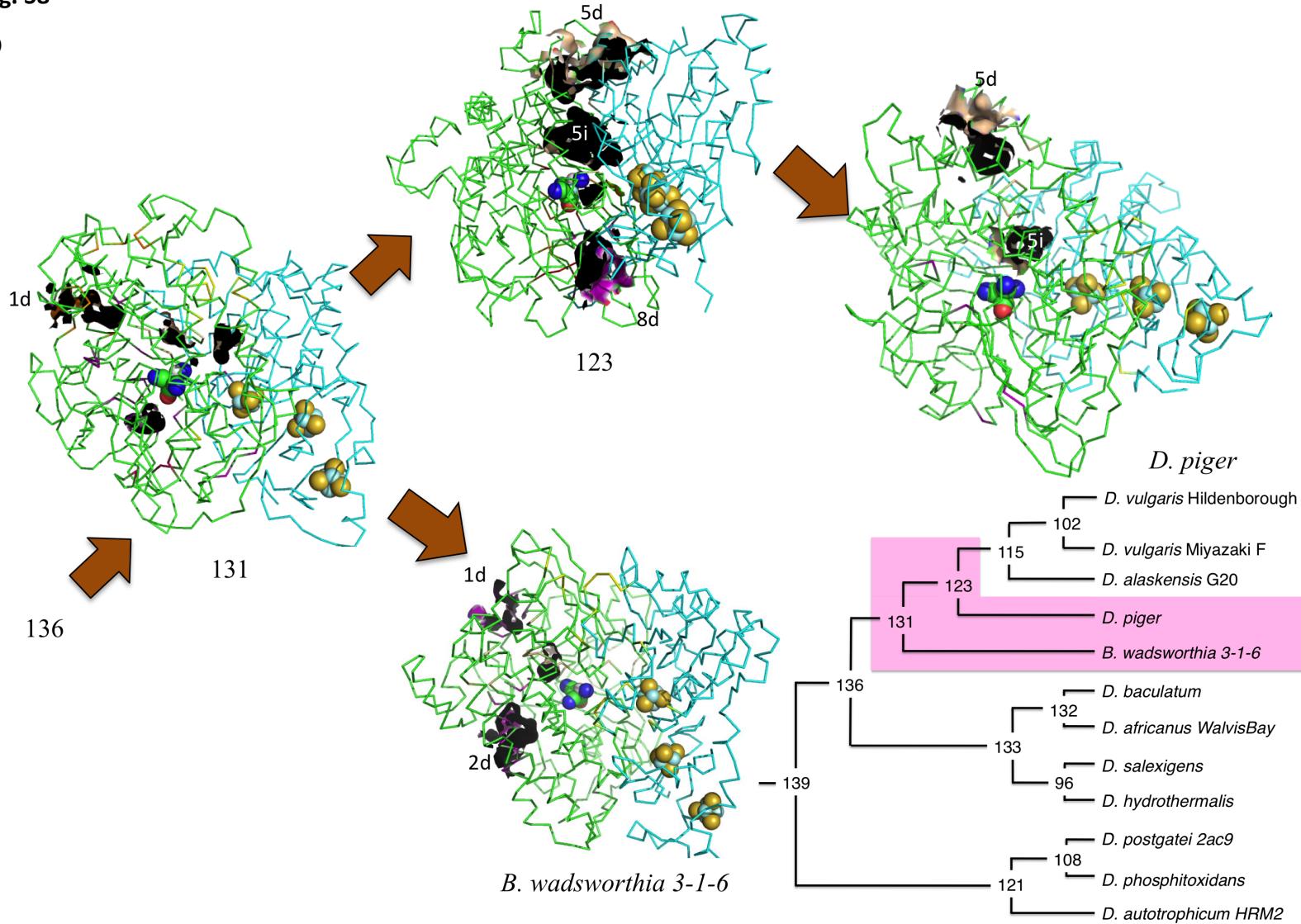


Fig. S8

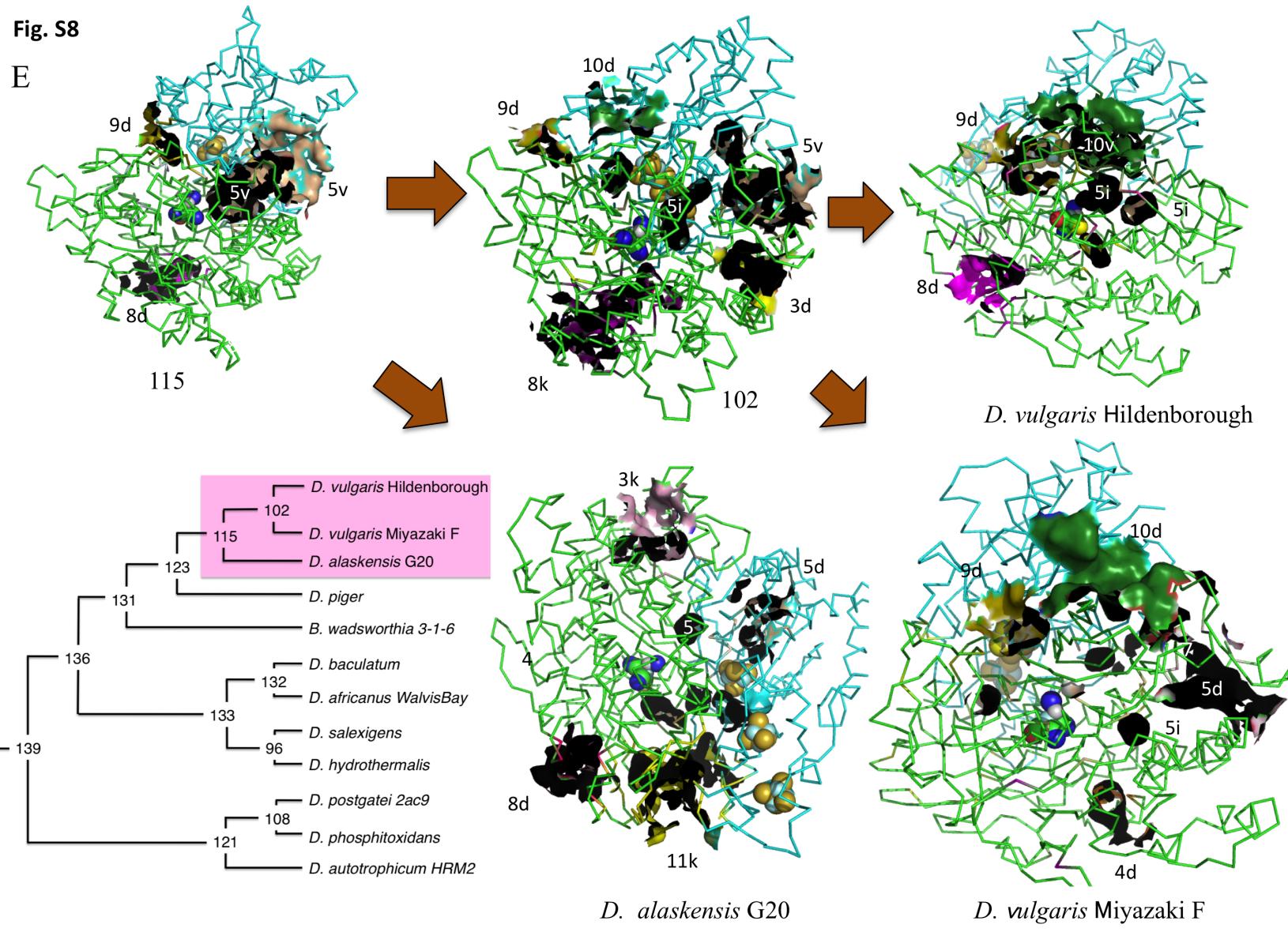
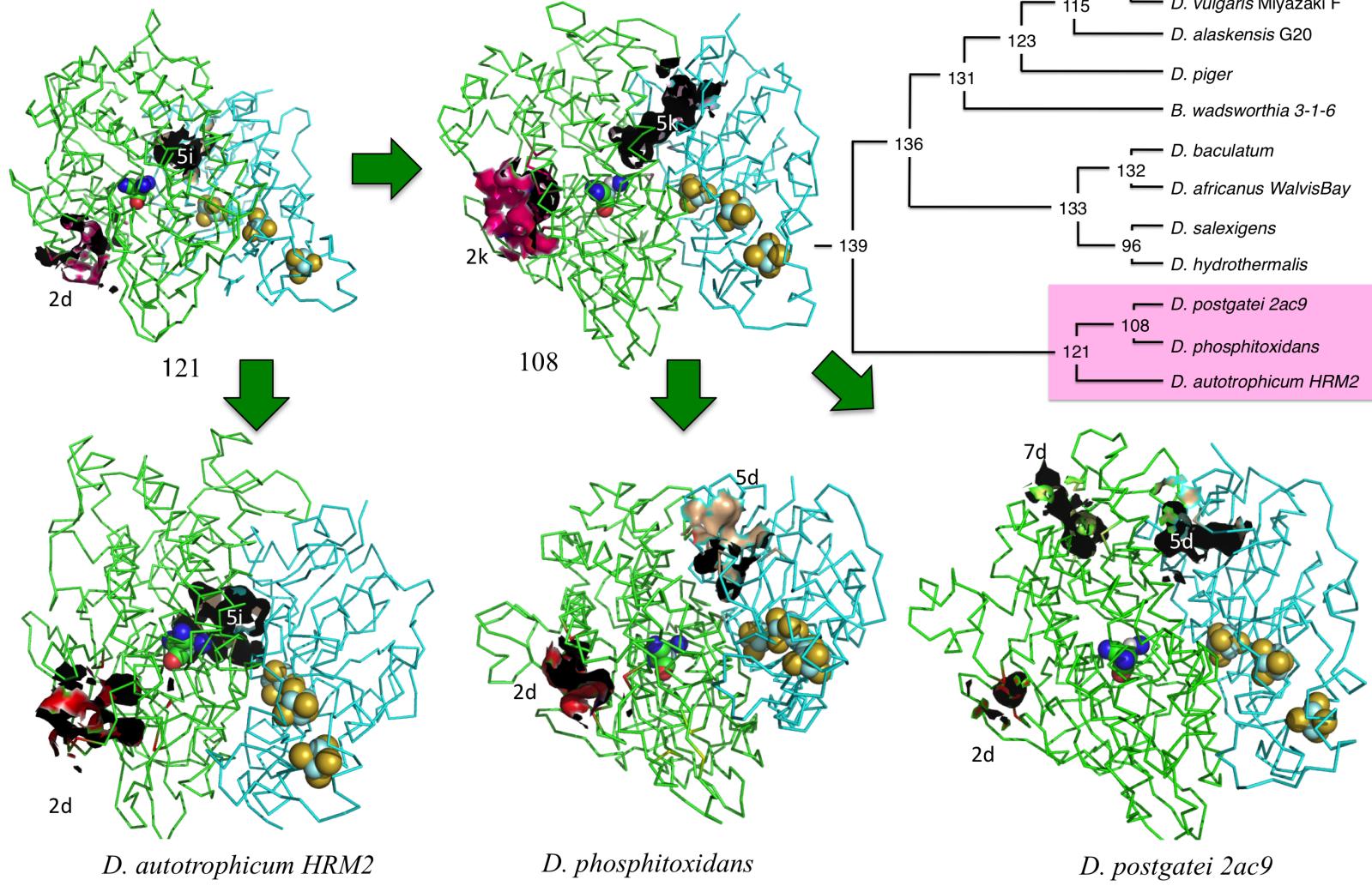


Fig. S8

F



Supplementary Figure S8: History of cavity development in [NiFeSe]Hases Comparative morphology of the cavities in [NiFeSe] Hases from the common ancestor 136 to the extant [NiFeSe]Hases *via* the intermediate ancestral forms. Cavities are identified by location and amino acid residues involved, and are designated by numbers and morphological designations, d, k, v, and i. The Ni-Fe and Fe-S clusters are shown in sphere designation, and the large and small subunits are designated in green and blue wire forms, respectively. (A-C) Blue arrows show evolutionary trait under absolutely anaerobic aquatic environments. (D, E) Brown arrows show path to terrestrial divergence including human faeces-derived isolates. (F) Green arrows show the path to marine sediment-derived [NiFeSe] Hases. Phylogenetic tree on each panel A-F highlights the genetic trait where the panel illustrates.

Supplementary Fig. S9

supplementary Fig. S9

Fig. S9

<i>D. vulgaris</i> Hilden. 102 <i>D. vulgaris</i> Miyazaki F 115 <i>D. alaskensis</i> G20 123 <i>D. piger</i> 131 <i>B. wadsworthia</i> 3-1-6 136 133 132 <i>D. baculatum</i> <i>D. africanus</i> WalvisBay 96 <i>D. hydrothermalis</i> <i>D. salexigens</i> 121 <i>D. autotrophicum</i> HRM2 <i>D. postgatei</i> 2ac9 108 <i>D. phosphitoxidans</i> 139	250 260 270 280 290 300 PDWIVGTVVLA-----LDAIKKNGL-----EGGLAE----VVKVLDSGRPTPFF PDWIVGTVVVA-----LNAIKENGL-----EAGLAE----VVKLLDADGRPTP-Y PDWIVGTVVVA-----LNAIKAKGL-----AAGLGD----VVKLLDADGRPTPFY PDWIVGTVVVA-----LDAIKENGL-----EAGLAE----VVKLLDADGRPTP-Y PDWIVGTVVVA-----LDAIKKHGL-----QGGLGE----VVKLLDGDRPTPFY PDWIVGTVVVA-----LNAIKENGL-----EAGLAE----VVKLLDADGRPMFY PDWIVGTVVVA-----LQAIKEKGL-----EAGLAE----VVSLLDSEGRPLPFY PDWIVGTVVVA-----LNAIKENGL-----EAGLAE----VVKLLDADGRPMFY- PDWIVGTVIGL-----LQALATNTL-----GLL-----VKQQLDANGRPAFY PDWIVGTLVVA-----LNAIKENGL-----EDGLAE----VVKILDDGRPMLF- PDWMVGTIVVA-----LNAIKENGL-----EDGLAE----VVKILDDGRPMLF- PDWMVGTIVVA-----LNAI-----L-----EHGLAE----VVKILDDGRPMLF- PDWMVGTIVAA-----WSHVLNPTEHPLPE-----LDDDRPMLFF PDWIIGSLAFA-----L-----EHGVREATAKI----LDAEGRPSVFF PDWMVGTIVVAINAIEEKG-----L-----QGGLAEVVKI----LDDDRPTP-F PDWMVGTIVVAINAIEEKG-----L-----QGGLAEIVTI----LDDEGRPTPFF PDWMVGTIVVAINAIEAKG-----L-----QGGLAEVVKI----LDDNGRPTPFF PDWIVGTVIHL-----L-----KEGLPE----LDDDRPMLFF PDWIVGTAIAHL-----L-----TKGLPE----LDENGRPMLFF PDWI VLS I V H-----L-----EKGIPE----LDDEGRPMLFF PDWIVGSIVH-----L-----KEGIPE----LDDDRPMLFF PDWIVGSIVH-----L-----NAGIPE----LDYDGRPTLFF PDWIVGTLV-----HVL-----EDGLPE----LDDDRPMLF-
<i>D. vulgaris</i> Hilden. 102 <i>D. vulgaris</i> Miyazaki F 115 <i>D. alaskensis</i> G20 123 <i>D. piger</i> 131 <i>B. wadsworthia</i> 3-1-6 136 133 132 <i>D. baculatum</i> <i>D. africanus</i> WalvisBay 96 <i>D. hydrothermalis</i> <i>D. salexigens</i> 121 <i>D. autotrophicum</i> HRM2 <i>D. postgatei</i> 2ac9 108 <i>D. phosphitoxidans</i> 139	310 320 330 340 350 360 GRNIHENCPYLDKYDEGVMSATFTDKVGCRYDLGCKGPMTADCFCERKWNGGVNWCVQNA GRNIHENCPYLDKYDEGKMSETFT-KDGCRYD-GCKGPMTMSDCFCERKWNGGVNWCVHNA GRNVHENCPYLEADAGKMCETFTKKEGCRYDLGCKGPMSMCDSFERKWNGGVNWCIENA GRNIHENCPYLDKFDEGMSETFT-KDGCRYD-GCKGPMTMSDCFCERKWNSGVNWCVDNA GTNIHDNCPYLPQFEYYVMSET TFT QKDGCRYELGCKGPSTMADCYKRKWNNNGVNWCISNA GRNIHENCPYLDKFDEKLSETFT-KDGCRYD-GCKGPATNSDCFCERKWNSGVNWCVDNA GRNVHENCPYLGYDEGKFSATFTEDGCRYDLGCKGPAYCDSFERKWN-GVNWCVANA QONIHENC PYLDKFDEGKLAETFT -KDGCRYD-GCKGPATNSDCFCERKWNSGVNWCVDNA -KNVHMNCPHLSAFEAGHMVKTMSDKDGCRFSMGCKGP SACDSFERKWNNNGVNWCVNNNA GENIHDNC PYLDKFNDKFAETFT -KDGCRYD-GCKGPATNSDCFCERKWNSGVNWCVDNA GENIHDNC PYLDKFNDKFAETFT -KAGCKYD-GCKGPATNSDCFCERKWNSGVNWCVENA GENIHDNC PYLDKYDNSEFAETFT KP-GCKAELGCKGP STYADCAKRRWNNGINWCVENA GENIHDNC PYLEHFENDNFAATFTQAG -CKYNLGCKGP ACNSDCFCERKWNSGMNWCVENA GENIHDNC PYLEAFDNDEYAEIFT -PDKCRYE-GCKGP SANSDCFKRKWNNGVNWCVENS GENIHDNC PYLEAFDNDEYAEIFT TDPEKCRYELGCKGP SANSDCFCERKWNSGVNWCVENS GENIHDNC PYLEAFDNDEYAEIFT DPVKCRYELGCKGP SANSDCFCERKWNSGVNWCVENS GENIHDNC PYLDDFDEDIFAEFT TNDKGCRMDLGCKGP DTYADCFKRWNSGLNWCVDNA GENIHDNC PYLDYFDQDIYSKTFTDKKGCRMEL GCKGP DTYADCFRRKWNNSGLNWCIENA GENIHDNC PRLKMYEADQLSQLTS SDPKGCRN L GCKGP STYADCYKRKWNSGLNWCVDNA GENIHDNC PYLDDYDAFMAATL SDPKGCRMDLGCKGP DTYADCYQRKWNSGLNWCVNA GENIHDNC PYLDKFNDKFAETFT -KDGCRFD-GCKGP ATYADCFKRWNSGVNWCVDNA

supplementary Fig. S9

Fig. S9

D. vulgaris Hilden.
102
D. vulgaris Miyazaki F
115
D. alaskensis G20
123
D. piger
131
B. wadsworthia 3-1-6
136
133
132
D. baculatum
D. africanus Walvis Bay
96
D. hydrothermalis
D. salexigens
121
D. autotrophicum HRM2
D. postgatei 2ac9
108
D. phosphitoxidans
139

370	380	390	400	410	420
....					
VCIGCVERPDFDGKSPFYQA—MSGCTPK—AAPA—GAT-----				GRTTIAIDPVTRIE	
VCIGCVERPDFPD—KSPFYQ—MSGCTPK—AAPA—GAT-----				GKTTIAIDPVTRIE	
VCIGCVERPDFDGKSPFYSA—MSGCTPK—AAPA—GAT-----				GKATIAIDPVSRIE	
VCIGCVERPDFPD—KSPFYE—MSGCTPK—TAPA—GAT-----				GKTTIAIDPVTRIE	
VCIGCVERPDFDGKSPFYES—MSGCTNK—MAAG—GVS-----				GKTKIAIDPVTRIE	
VCIGCVERPDFPD—KSPFYEM—T—TPK—TAPA—GAN-----				GKTTIAIDPVTRIE	
IICIGCTEPSPFDGQSPFYSN—MA-----				KATIAIDPVTRIE	E
VCIGCVERPDFPD—KSPFYEM—T—TSK—TAPA—GAN-----				GKTTIAIDPVTRIE	
TClGCTSPTFPDQGQSPFYVN—MS-----				KTVIAIDPVTRLE	
VCIGCVERPGFPD—MSPFYEM—T—ASK—TAPA—GAD-----				GKIKIAIDPVTRIE	
VCIGCVERPGFPD—MSPFYEM—M—ASK—TAPA—GAD-----				GKIKIAIDPVTRIE	
VCIGCVERPGFPD—MSPFYEM—M—ASK—TAPA—GAD-----				GKVKISIDPVTRIE	
VCIGCVERPDFDGKSPFYVAE—VS—Q—AATP—AAD-----				GKVKISIDPLTRVE	
VCTGCAEPGWPDNFSPFYESM—MS—EKN-----				VKISIDPVTRIE	
VCIGCVERPGFPD—MSPFYE—G—MS—SKS—HAPA—GKD-----				GKIKIAIDPVTRIE	
VCLGCVEPGFPDEMSPFYEAG—MS—SKS—HAPA—SKD-----				GKIKIAIDPVTRIE	
VCIGCVERPGFPDEMSPFYEAG—MS—SKS—HAPA—GKD-----				GKIKIAIDPVTRIE	
VCIGCVERPGFPDAMSPFYEPA—MAGCKPE—AAPAGATG-----				KKIKIAIDPVTRIE	
VCIGCVERPGFPDASSPFYEQS—MAGCKPE—AVPVGAA-----				KKIKVAIDPVTRIE	
VCIACVERPGFPDQSSPFYEPA—MSGSKP-----			TTGSVGV DSTKKLKISIDPI	TRIE	
VCIGCVERPGFPDAMSPFYEPA—MAGCKPQ—AEPA—ATG-----				KKIKIGIDPVTRIE	
VCIGCVERAGFPDAMSPFYEPAYMAGCQPQ—AEPA—QTG-----				KKIKIGIDPVTRIE	
VCIGCVERPGFPD—MSPFYEM—T—ACKPATAPA—GAD-----				GKIKIAIDPVTRIE	

D. vulgaris Hilden. 102
D. vulgaris Miyazaki F 115
D. alaskensis G20 123
D. piger 131
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D. africanus Walvis Bay 96
D. hydrothermalis
D. salexigens 121
D. autotrophicum HRM2
D. postgatei 2ac9 108
D. phosphitoxidans 139

430 440 450 460 470 480
|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
GHLKAEVVENGKVVDARLSGGMYRGFETILGRGRDPRDASQIVQRI CGVCPТАHSTASVL
 GHLKAEVVENG-VVDARLSGGMYRGFETI LGRGRDPR-ASQIVQRI CGVCPТАHSTASVL
 GHLKAEVTENGVVVDARLSGGMYRGFETI LGRGRDPRDASQIVQRI CGVCPТАHSTASCM
GHLKAEVVENG-VVDARLSGGMFRGFENILRGRDPR-ASQIVQRI CGVCPТАHSTASVL
 GHLKAEVVVEGGKVVDAHISGGMFRGFENIL RGRDPRDASQIVQRI CGVCPТАHSTASVM
 GHLKAEVVVEDG-VVDARLSGGMFRGFENIL SGRDPR-ASQIVQRI CGVCPТАHATASAL
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 GHLKAEVEVEDG-VVDARLSGGMFRGFENIL SGRDPR-ASQIVQRI CGVCPТАHATASAL
 GHLKVEVQVEDGKVADAWI TGGMFRGFEI LGRGRNPRDASQIVQRI CGVCPVAHATASSL
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GHLKAEVEVEDG-VVDWFSGGMFRGFENILIGRDPR-ASQIVQRI CGVCPТАHSTASAL
GHLKIEVEVKDGKVVDAKCSGGMFRGFENILIGRDPRDS SQIVQRI CGVCPТАHCTASVM
GHLKAEVKENGVVTDAMSSGGMFRGFENILIGRDPRDATQIVQRI CGVCPТАHSTASCL
 GHLKAEVVVKDG-VVDWFSGGMFRGFENIL IGRDPR-AAQLTQRLCGVСRTAHTASTL
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 GHLKAEVVVKDGKVTDWLSSGGMYRGFENIL IGRDPRDAAQLTQRLCGVCPТАHSTASR
GHLKAEVEVEDGKVVDARFSGGMFRGFENILNGRDPRDATQIVQRI CGVCPТАHATASAL
 GHLKVEVEVKGKVVNDARCFGGMFRGFENIL TGRDPRDATQIVQRI CGVCPТАHATASSL
 GHLKAEVEVKNGVVVDARMSGGMYRGFEEQIL VGRDPRDAVQITQRI CGVCPТАHATASSL
 GHLKAEVEVEDGKVVDARISGGMYRGFEEQIL NGSDPRDATQITQRI CGVCPТАHATASAL
 GHLKAEVEVAGGKVVDAHITGGMYRGFEEQIL YGRDPRDATQITQRI CGVCPТАHATASAL
 GHLKAEVEVEDG-VVDWFSGGMFRGFENIL IGRDPR-AAQIVQRI CGVCPТАHATASAL

supplementary Fig. S9

	490	500	510	520	530	540
<i>D. vulgaris</i> Hilden.	ALDEAFGAKVPNNGRITRNLI	F	GANYLQSHILHFYHLSAQDFVQGPDTAPFVPRFPKSDL			
102	ALDD-FGVKPTNGRI-RNLIF		GANYLQSHILHFYHAA	L-FVQGPDTAPFV-RFAKPDL		
<i>D. vulgaris</i> Miyazaki F	ALDN	AFKVVKPTNGRLTRNL	TFGANYLQSHILHFYHAA	LDL-FVQGPDTAPFV-RFAKPDL		
115	ALDD-FGVKVTTNGRI-RNLIL	GANYLQSHILHFYHAA	L-FVQGPDTAPFV-RFAKPDL			
<i>D. alaskensis</i> G20	ALDKA	FGVKVTTNGRLTRNL	LGANYMQSHILHFYHAA	ALDFVQGPETAPFVPRFKNPDL		
123	ALDD-FGVKVTTNGRI-RNLIL	GANYLQSHILHFYHAA	L-FVQGPDTAPFV-RFAKPDL			
<i>D. piger</i>	AQE	DAFNIVKTGNGRITRNLI	LGANYLQSHILHFYHAA	LDL-FVAGPDTAPFVPRFAQPDL		
131	ALDD-FGVKVTTNGRI-RNLIL	GANYLQSHILHFYHAA	L-YVQGPDTAPFV-RYDNPDL			
<i>B. wadsworthia</i> 3-1-6	AIEAVCGVEVPENGRIARNLMLAGNYLQSNILHFYHLLGGQDYFHGPD	QDYFHGPD	TVFIPRYRNPD			
136	ALDD-FGVKVTTNGRI-RNLIL	GANYLQSHILHFYHAA	L-YVNGPDMAPFV-RYDNPDL			
133	ALDD-FGVKVTTNGRI-RNLIF	GANYLQSHILHFYHAA	L-YVNGPDIAPFV-RYDNPDL			
132	ALDD-FGVKVTTNGRI-RNLIF	GANYLQSHILHFYHAA	L-YVNGPDIAPFV-RYDNPDL			
<i>D. baculum</i>	AODDA	FGVKVTTNGRITRNLI	FGANYLQSHILHFYHAA	LDYVKGPDVSPFVPRYANADL		
<i>D. africanus</i> WalvisBay	ALDDA	FGVKLT	TTNGRVT	FGANYLQSHILHFYHAA	LDYVAGPDVAPFVPRYEKSDL	
96	ALDD-FGVKL	TTNGRVT	RNLIF	FGANYLQSHILHFYHAA	L-YVNGPDIAPFV-RYDNPDL	
<i>D. hydrothermalis</i>	ALDDAFGAK	ITTNGRVTRNL	IFGANYLQSHILHFYHAA	L-TALDFVRGPGKAPFVPRFEQPD		
<i>D. salexigens</i>	ALDDAFG	VKLT	NNGRVTRNL	IFGANYLQSHILHFYHAA	LDVFRGPKAPFVPRFEQPD	
121	ALDDAFG	VKLT	NNGRVARNL	ILGANFLQSHILHFYHAA	LDVFRGPKAPFVPRFEQPD	
<i>D. autotrophicum</i> HRM2	ALDDA	FGVKLT	NNGRVARNL	ILGANFLQSHILHFYHAA	LDVFRGPKAPFVPRFEQPD	
<i>D. postgatei</i> 2ac9	ALDDAFG	VTL	TDN	GRVARNL	ILGANFLQSHILHFYHAA	LDVFRGPKAPFVPRFEQPD
108	ALDDAFG	VKL	TDN	GRVARNL	ILGANFLQSHILHFYHAA	LDVFRGPKAPFVPRFEQPD
<i>D. phosphitoxidans</i>	ALDDAFG	VEL	TDN	GRVARNL	ILGANFLQSHILHFYHAA	LDVFRGPKAPFVPRFEQPD
139	ALDD-FGVKV	TNNGR	IARNL	IFGANYLQSHILHFYHAA	L-YVNGPDMAPFV-RYDNPDL	
	550	560	570	580	590	600
<i>D. vulgaris</i> Hilden.	RLSKELNKAG	-----	VDQYIEALEVRRICHEMVALFGGRMPHVQGQVVGGATEIPT			
102	RLPKEMNKA	-----	VDQYLEALEVRRICHEMVAL-GGGRMPHVQGQVVGGATEIPT			
<i>D. vulgaris</i> Miyazaki F	RLPKDINAAA	-----	VDQYLEALEVRRICHEMVALFGGRMPHVQGQVVGGATEIPT			
115	RLPKEMNKV	-----	VDQYLEALEVRRICHEMVAL-GGGRMPHVQGQVVGGATEIPT			
<i>D. alaskensis</i> G20	RLPSAVNQVA	-----	VDQYLEALEVRRICHEMVAIFGGRMPHVQGQVVGGATEIPT			
123	RLPKEMNKV	-----	VDQYLEALEVRRICHEMVAL-GGGRMPHVQGQVVGGATEIPT			
<i>D. piger</i>	RLPPEANKVG	-----	VDQYLEALEVRRIAHEMVALFGGRMPHVQGIVPGGATEMPT			
131	RLPKEMNKV	-----	VDQYLEALEVRRICHEMVAL-GGGRMPHVQGQVVGGATEIPT			
<i>B. wadsworthia</i> 3-1-6	RLSEEQNTLA	-----	MDEYIEALEVRQVCHQLVALFGGRMPHLQGILGGGAAQIPD			
136	RLAKEINKV	-----	VDQYLEALEIRRICHEMVAL-GGKMPHVQGMVVGGATEIPT			
133	RLAHEINKV	-----	VDQYLEALEIRRICHEMVAL-GGKMPHVQGMVVGGATEIPT			
132	RLAHEINKV	-----	VDQYLEALEIRRICHEMVAL-GGKMPHVQGMVVGGATEIPT			
<i>D. baculum</i>	-LTDRIKDGAKADNTYGLNQYLKALE	IRRICHEMVA	FGGRMPHVQGMVVGGATEIPT			
<i>D. africanus</i> WalvisBay	RLTPELNKVA	-----	VDQYLEALNVRVLVAHEMVALFGGKMPHVSGQVVGGTEIPS			
96	RLDEKTNKV	-----	VDQYVKALEIRRICHEMVAL-GGKMPHVQSGQVVGGATEIPT			
<i>D. hydrothermalis</i>	RLDEK	TNAVA	-----	VDQYIKALEIRRICHEMVALFGGKMPHISGQVVGGTEIPT		
<i>D. salexigens</i>	RLDEKTNKVA	-----	VDQYVKALEIRRICHEMVALFGGKMPHVSGQVVGGATEIPT			
121	RLPKEINEVA	-----	VDQYLEALEIRKICHEMVALFGGKMPHVQGIIVGGTEIPT			
<i>D. autotrophicum</i> HRM2	RLDKATNQVG	-----	VDQYLEALEIRKICHEMVALGGKMPHVQGIIVGGTEIPT			
<i>D. postgatei</i> 2ac9	RVSKDINDLC	-----	VGQYLEALEIRKICHEMVALGGKMPHVQGIIVGGTEIPT			
108	RLPKEINDVA	-----	VGQYLEALEIRKICHEMVALGGKMPHVQGIIVGGTEIPT			
<i>D. phosphitoxidans</i>	RLPKELNDVA	-----	VGQYLEALEIRKICHEMVALGGKMPHVQGIIVGGSTEIPT			
139	RLAKEINKV	-----	VDQYLEALEIRRICHEMVAL-GGKMPHVQGMVVGGATEIPT			

supplementary Fig. S9

	610	620	630	640	650	660		
<i>D. vulgaris</i> Hilden.	KEKLVEYAARFKKVRDFVEQKYVPVYTI	GSKYKDMFKVGQGFKAAL	CVGAFPLDNGKK				
102		KEA-VEYAARFKKVREFVEE	KYVPV-YT	GTVKDLFKE	FGQGYKN	CISFGAFLNDDGNE		
<i>D. vulgaris</i> Miyazaki F	KEALVEYAARFKKVRKFVEEKYVPV	YI	GSVYKDLFA	FGQGYKN	CISFGVFLNDDGKE			
115		KEA-VEYAARFKKVRKFVEEKYVPV	YI	GSVYKDLF	FGQGYKN	CISFGVFLNDDGKE		
<i>D. alaskensis</i> G20	KEALVEYAARFKKVRKFVEEKYVPV	YI	GSVYKDLF	FGQGYKN	CISFGVFLNDDGKE			
123		KEA-VEYAARFKKVRKFVEEKYVPV	YI	GSVYKDLF	FGQGYKN	CISFGVFLNDDGKE		
<i>D. piger</i>	KEALLEYAARFKKVRQFIVEKYLPI	TYI	GSVYKDLF	FGQGYKN	CISFGVFLNDDGKE			
131		KET-VEYAARFKKVRKFVEEKYVPV	YI	GSVYKDLF	KI	FGQGYKN	CISFGVFLNDDGKE	
<i>B. wadsworthia</i> 3-1-6	RETILEYAARMKQVRKFVENRYLPL	VYT	ASRYMDM	FEMAHG	GYKNA	LCAVGVFPLAKKG		
136		KEK-AEYAARFKKVRKFIEEKYVPV	-YI	MSVYTDLF	KI	GEQYKN	IAFGVFLNDDGKE	
133		KEK-AEYAARFKKVKHFIEEKYVPV	-YI	MSVYTDLF	KI	GDGYKN	IAFGVPMDDGKE	
132		KEK-AEYAARFKKVKHFIEEKYVPV	-YI	MSVYTDLF	KI	GDGYKN	IAFGVFPMDDGKE	
<i>D. baculatum</i>	ADKVAEYAARFKEVQKFVIEEYL	P	IYTLGSVY	TDLF	E	GIGWKN	IAFGVFPEDDYKT	
<i>D. africanus</i> WalvisBay	QQKLD	EYTKR	FKLVRKF	IEETYV	PVY	TKAYADLLK	VGDGYKN	CISFGVFPMDSGKE
96		KEK-AEYASRFKEVQKFIEE	TYVP	-VYI	GSVYKDLF	KI	GGGGYKN	AMAYGVFPMDAGYE
<i>D. hydrothermalis</i>	KEKLAEYASRFKDQVQKF	IEE	TYV	I	GSVYKDLF	KI	GGGGYKN	CISFGVFPMSDGDD
<i>D. salexigens</i>	KEKLAEYASRFKDQVQKF	IEE	TYV	I	GSVYKDLF	KI	GGGGYKN	AMAYGVFPMDAAESE
121		RENLDAYAERFKKVRKFIEEKY	VPIVY	LLLAGP	YGDLLK	TGVG	YKN	IAFGVFPPLDEGN-
<i>D. autotrophicum</i> HRM2	REKLDAYKERFKTVRKFIEERY	YLPL	IYLLAGP	YGDLLK	TGTG	YKN	IAFGVFPPLDEGN-	
<i>D. postgatei</i> 2ac9	REALNAYAERFKKIKKFVMEK	YI	PVYI	LAGPY	GDLLK	TGVGHK	NLVSWG	VFPFLDNKGN-
108		REALNAYAERFKKVRKFIVEK	YVPIVY	LLAGP	YGDLLK	TGVGHK	NLVSWG	VFPFLDNKGN-
<i>D. phosphitoxidans</i>	REALNAYAERFKKVRQFILEK	YVPIVY	LLAGP	YGDLLK	TGVGHK	NLVSWG	VFPFLDNKGN-	
139		KEK-AEYAARFKVSKFIEEKY	VPV	-YI	MSVYTDLF	KI	GEQYKN	IAFGVFPEDDGNE
	670	680	690	700	710	720		
<i>D. vulgaris</i> Hilden.	HLFMPGVYAKGKDMPFDPS	KI	KEYVKYSW	FAEETTG-LNYKEGKT	IPAPDKAGA	YSFVKA		
102	LLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-KEGKT	IPAPDKA	-AYSFVKA
<i>D. vulgaris</i> Miyazaki F	LHLKR	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-KEGKT	IPAPDKA	-AYSFVKA
115	LLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-NEGKT	IPDPDKA	-AYSFVKA
<i>D. alaskensis</i> G20	FALKPGVYMDGEDKPF	DARL	KEYVKYSW	-DDATTG-LH	-TEGQT	IPDPDKA	-AYSFVKA	
123	LLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-NEGKT	IPDPDKA	-AYSFVKA
<i>D. piger</i>	LLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-NEGKT	IPDPDKA	-AYSFVKA
131		HLLKAGVFLNGRD	VEFDP	KKITEDL	KYAWYD	ATTG-KGADGA	ETNPNL	DKKDAYSFVKA
<i>B. wadsworthia</i> 3-1-6	LLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-NEGKT	IPDPDKA	-AYSFVKA
136	QFFNAGAYINGRDEPFDG	NRILED	VRYSWFEPAPSG	-TPLQKS	ESNPQVD	KEGAYSF	FIKA	
133	FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDATTG-LH	-SEGKT	IPDPDKA	-AYSFVKA
132	FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKA	-AYSFVKA
<i>D. baculatum</i>	FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKA	-AYSFVKA
<i>D. africanus</i> WalvisBay	FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKP	-AYSFVKA
96		TLLKP	GVYIDGKDVAFDPA	KI	KEYAKYSW	FEDSCSN-LHPSQ	GKTLPKLG	PGAYSF
<i>D. hydrothermalis</i>	FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKA	-AYSFVKA
<i>D. salexigens</i>	FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKP	-AYSFVKA
121		FLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKA
<i>D. autotrophicum</i> HRM2	TLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKA	-AYSFVKA
<i>D. postgatei</i> 2ac9	TLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKP	-AYSFVKA
108		TLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKA
<i>D. phosphitoxidans</i>	TLLKP	GVYIDGKDP	FPDKQ	KEYVKYSW	-DDASTG-LH	-SEGKT	IPDPDKP	-AYSFVKA
139		-LLKP	GVYIDGKDP	FPDNQ	KEYVKYSW	-DDDTG-LH	-SEGKT	IPDPDKA

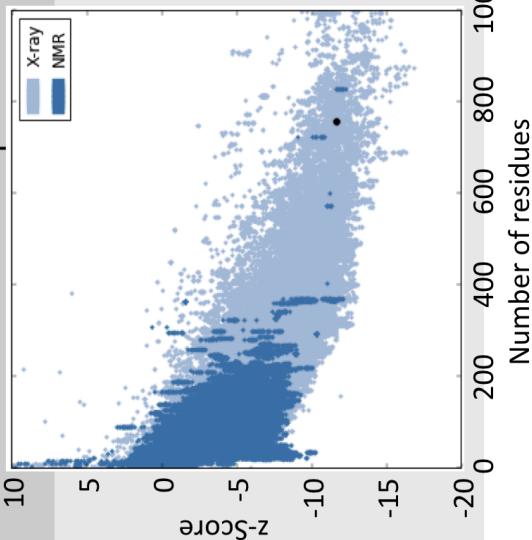
supplementary Fig. S9

	730	740	750	760	770	780
<i>D. vulgaris</i> Hilden.						
102	PRYDGLSLEV G PLARMVNNPELSPVGKLLKDLFGISAKKFRDLG-EAAFSLMGRHVA					
<i>D. vulgaris</i> Miyazaki F	PRYNKPVEVG P LA-MWVT-PELSPVGQKLLKDLFGIDAKSFRD-G-DEAAFS L MGRHVA					
115	PRYNKAVES G PVARMWI-NPELSPVGQKLLKDLFKLNAKRFRDLG-DEAAFSVMGRHVA					
<i>D. alaskensis</i> G20	PRYNKPVEVG P LA-MWVT-PELSPMGQKLLKDLFGIDAKSFRD-G-DEAAFSLMGRHVA					
123	PRYNKPVEVG P LA-MWVT-PELSPMGQKLLKDLFGIDAKSFRD-G-DEAAFSIMGRHVA					
<i>D. piger</i>	PRYDG E VIEVGPAAR M WVANAPLSEVGVKMLKEKFGIEARTIRDLG-WDKVFSIMGRHVA					
131	PRYNKPVEVG P LA-MWVT-PELSPMGQKLLKDLFGIDAKSFRD-G-DEAAFSIMGRHVA					
<i>B. wadsworthia</i> 3-1-6	PLYAGHR E V G PLARM W NDKPLSPIGQRFFADM F VG R VRAETFRQIG-EDPAFSIMGRNVA					
136	PRYNKPVEVG P LA-MWVT-PELSPMGQKLLKDLFGIDAKSFRD-G-DEAAFSIMGRHVA					
133	PRYNKPVEVG P LA-MWVP-PELSPMGQKLLKDLFGIDAKSFRD-G-DEMAFSIMGRHVA					
132	PRYNKPVEVG P LA-MWVP-PELSPMGQKLLKDLFGIDAKSFRD-G-DEMAFSIMGRHVA					
<i>D. baculatum</i>	PRYKD K PCEV G PLARM W VQNPELSPVGQKLLKELY G IEAKNFRDLG-DKAFSIMGRHVA					
<i>D. africanus</i> WalvisBay	PRYDGH P VEVG P LA-MWVP-PELSPMGKKQLKDLFGINANRFRDLG-DLAFS I MGRHVA					
96	SRYNK A VEVG P LA-MWVH-PELSPMGKKQLKDLFGIEAKIFRD-GE-DEMAFSLMGRHVA					
<i>D. hydrothermalis</i>	SRYNNAVEVG P LA-MWVH-PELSPMGKKQLKDLGY I TAKNFRDLGE-DEMAFSLMGRHVA					
<i>D. salexigens</i>	SRYNK A VEVG P LA-MWVH-PELSPMGKKQLKDLFGIEAKMFRDLGE-DEMAFSLMGRHVA					
121	PRYNKPHEVG P LA-MWVTPNELSAT G Q K AL-----GVKRLSDIG-DAAFSILGRHVA					
<i>D. autotrophicum</i> HRM2	PRYNKPHEVG P LA-MWVTPNELSAT G Q K AL-----GVKRM R DIG-DAAFSILGRHIA					
<i>D. postgatei</i> 2ac9	PRYNKPHEGG P LA-MWVTPNELSAT G Q E AL-----GVKKLRDIG-DACFSILGRHVA					
108	PRYNKPHEGG P LA-MWVTPNELSAT G Q K AL-----GVKRLSDIG-DACFSILGRHVA					
<i>D. phosphitoxidans</i>	PRYNKPHEGG P LA-MWVTPNELSAT G Q K EL-----GVTRLRDIG-DACFSILGRHVA					
139	PRYNKP L EV G PLA-MWVT-PELSPMGQ K LL-----DAKSFRD-G-DEAAFSIMGRHVA					
	790	800	810	820	830	840
<i>D. vulgaris</i> Hilden.						
102	RAEETYYMLGAIE--GWLKEIKAGE D TV V MPAVPASAEGTG F TEAP R GSLLHYVKVKDSK					
<i>D. vulgaris</i> Miyazaki F	RAEETYYMLNAIE--RWLKE V KAGEET-VASEIP-SAE G IG F TEAP R GSLLHYINIK-SK					
115	RAEETYY M SAIE--RWLKEV K AGEETFAAAEIPASSEG V G F TEAP R GSLLHYINIKDQK					
<i>D. alaskensis</i> G20	RAEETYLIPNAIE--RWLKEVK P GEET-VPSEIP-SAE G IG F TEAP R GSLLHYINIK-YK					
123	RAEETWLTNYIE--RWLKEV V PG A ET Y P S E I PEQAEGTGFTEAP R GSLLHY I D I KDSV					
<i>D. piger</i>	RAEETYLIANAIE--RWLKEV K PG E ET-VPSEIP-SAE G IG F TEAP R GSLLHYINIK-YK					
131	RAEALLIANA E --GWLKEV K PD G ET F TF P FE I P Q SAEG G C E AP R GS L VHYIRVKDQK					
<i>B. wadsworthia</i> 3-1-6	RAEETYLIANAIE--RWLKEV K PG E ET-VASEIP-TAE G IG F TEAP R GSLLHYINIK-YK					
136	RVEEVYQT L GM E --YWLHELEPG Q AT F ALPEV P Q A EG G IG F TEAP R GC L HYMRV K NG V					
133	RAEETYLVANAIE--RWLKEV K PG E ET-VASEIP-TAE G IG F TEAP R GS L HYINIK-YK					
132	RAEETYLVANA E D-RWLKEV K PG E ET-VT S EIP-TAE G IG F TEAP R GSLLHYINIK-YK					
<i>D. baculatum</i>	RAEETYLVANA E D-RWLKEV K PG E ET-VT S EIP-TAE G IG F TEAP R GSLLHYINIK-YK					
<i>D. africanus</i> WalvisBay	RAEETWLTA V AVE--KWLKV Q PG A ET Y K S E I P D AA E GT F TEAP R G A LLHY L K I K D KK					
96	RAEETYLVAKA E --ERWLKEV K PG K ET F V A AS I P D SAEG G IG F TEAP R GSLLHYVN I K D KK					
<i>D. hydrothermalis</i>	RAEEAYMV N AAIE--DAWL E K V PG E ET-VK T E I P-SAE G GL T EAP R GSLLHYINIK-SK					
<i>D. salexigens</i>	RAEESYLV N AA I G--DIWL E VG E GE E TYVK T MP E SG E VG V LT E AP R GSLLHYINIKDSK					
121	RAEEAYMV N AI Q --DAWL E Q V PG E ET Y K T E I P S AE G GL T EAP R GSLLHYINIKDSK					
<i>D. autotrophicum</i> HRM2	RAEETLLV A K M E--HWLDEAK P GE E T V AA I P N AE G IG L TEAP R G A LLHY I D I K N Y K					
<i>D. postgatei</i> 2ac9	RAEETLLV A M Q ME--RWL E EA K PG L ET F V A AP I P N AE G IG L TEAP R G A LLHY I D I K N SV					
108	RAEETLLV A K A E--QWIAQ A T P KG K ET F V P AA I P N AE G IG L GM T TEAP R G A LLHY I D I K N Y K					
<i>D. phosphitoxidans</i>	RAEETALV A A M E--EWL T Q A QP G K E T F V P AP I P D SAQ G LG M TEAP R G A LLHY I D I K D Q K					
139	RAEETYLVANAIE--HWLKEV K PG E ET-VASEIP-NAEG G IG F TEAP R G A LLHYINIK-YK					

supplementary Fig. S9

Supplementary Fig. S9: Amino acid sequence alignment and identification of residues involved in the cavity. Amino acid residues that have more non-synonymous substitutions ($\delta N - \delta S > 0$) are marked in yellow. The residues constituting cavities are designated in red. Assemblage names in the left column are grouped as terrestrial origin (highlighted in yellow), clinical isolates (brown letters), deep-sea marine vent origin (highlighted in blue), and sediment origin (highlighted in green). Ternary structural models start with R78 and end at H900.

MD model	z-Score	MD model	z-Score	MD model	z-Score
<i>D. vulgaris</i> Hildenborough 2wpn.pdb (<i>D. vulgaris</i> H.)	-11.71	<i>D. baculum</i>	-11.22	<i>D. postgatei</i> 2ac9	-9.86
<i>D. vulgaris</i> Miyazaki F	-11.59	1cc1.pdb (<i>D. baculum</i>)	-10.64	<i>D. phosphit-</i> <i>oxidans</i>	-9.99
<i>D. alaskensis</i> G20	-10.69	<i>D. africanus</i> Walvis Bay	-11.20	<i>D. autotrophicum</i> HRM2	-11.22
<i>B. wadsworthia</i> 3-1-6	-11.38	<i>D. salexigens</i>	-10.73	108	-11.10
<i>D. piger</i> 102	-10.42	<i>D. hydrothermalis</i>	-8.96	121	-11.25
115	-11.56	132	-9.69		
123	-10.25	96	-10.30		
131	-10.07	133	-9.92		
		-9.71	136	-9.75	
		-9.88	139	-10.42	



Supplementary Table S1: Quality of MD models evaluated by ProSA. ProSA is a knowledge-based program for the evaluation of the model accuracy. The z-score indicates overall model quality and measures the deviation of the total energy of the structure, which is displayed in a plot that contains the z-scores of all experimentally determined protein chains in current PDB. The range of z-scores between -9 and -14 is acceptable for the [NiFeSe]-Hases with 770 amino acid residue. X-ray crystallographic models 2wpn.pdb and 1cc1.pdb are also evaluated as reference.