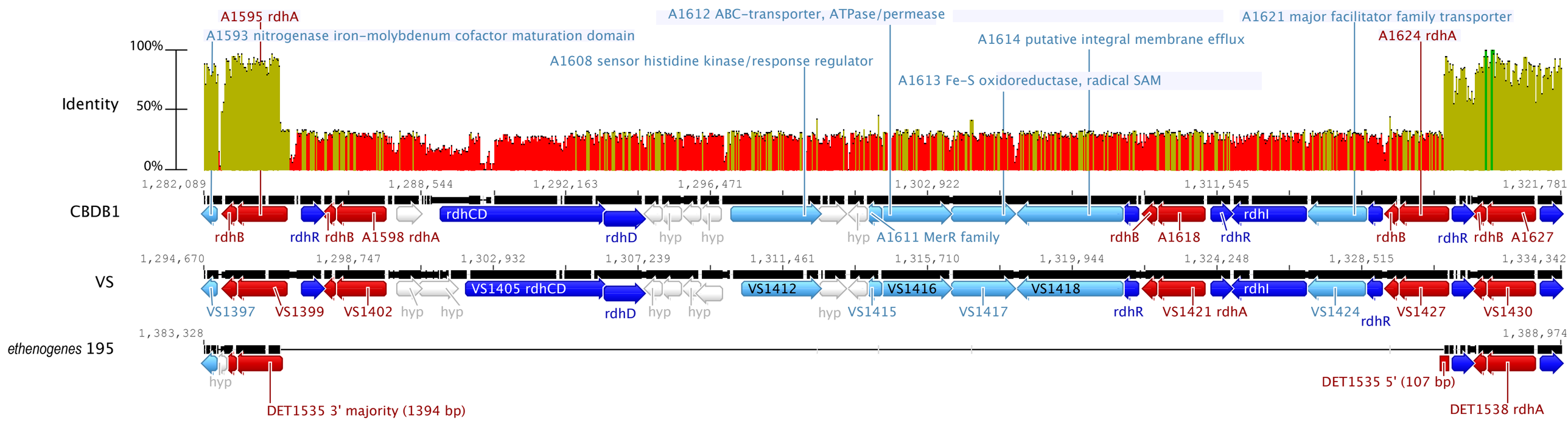
A1595 rdhA
A1593 nitrogenase iron-molybdenum cofactor maturation domain
A1612 ABC-transporter, ATPase/permease
A1621 major facilitator family transporter
A1624 rdhA
A1614 putative integral membrane efflux
A1608 sensor histidine kinase/response regulator
A1613 Fe-S oxidoreductase, radical SAM
100%
Identity
50%
0%
1,282,089
1,288,544
1,292,163
1,296,471
1,302,922
1,311,545
1,321,781
CBDB1
rdhB
rdhR
rdhB
A1598 rdhA
rdhCD
rdhD
hyp
hyp
hyp
A1611 MerR family
rdhB
A1618
rdhR
rdhI
rdhB
rdhR
rdhB
A1627
1,294,670
1,298,747
1,302,932
1,307,239
1,311,461
1,315,710
1,319,944
1,324,248
1,328,515
1,334,342
VS
VS1397
VS1399
VS1402
hyp
hyp
VS1405 rdhCD
rdhD
hyp
hyp
VS1412
hyp
VS1415
VS1416
VS1417
VS1418
rdhR
VS1421 rdhA
rdhI
VS1424
rdhR
VS1427
VS1430
1,383,328
1,388,974
ethenogenes 195
hyp
DET1535 3' majority (1394 bp)
DET1535 5' (107 bp)
DET1538 rdhA