

Dataset S1. Radius and hydrophobicity profiles for all ion channel structures analysed. Each channel structure is oriented with the cytoplasmic (or mitochondrial matrix) side in the negative s -axis direction. The approximate position of the lipid bilayer membrane, centred at $s = 0$ nm, is indicated by a beige shading. Simulation results are coloured red if the main barrier to water permeation has height > 2.6 kJ mol⁻¹ and blue otherwise, with a similar colour divide for the heuristic scores, at a cut-off of 0.55.

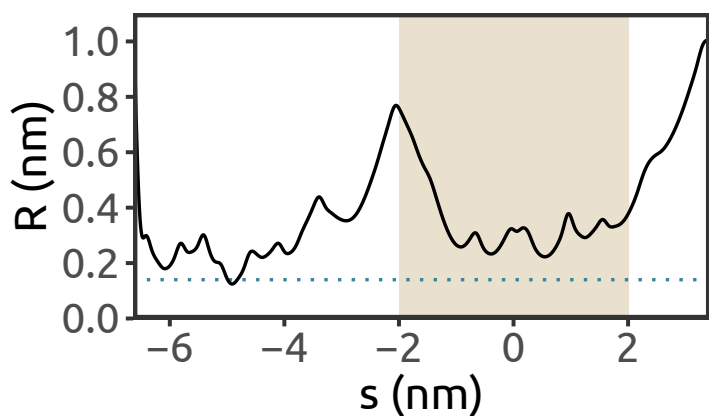
5HT3R (PDB ID: 4PIR)

Mus musculus

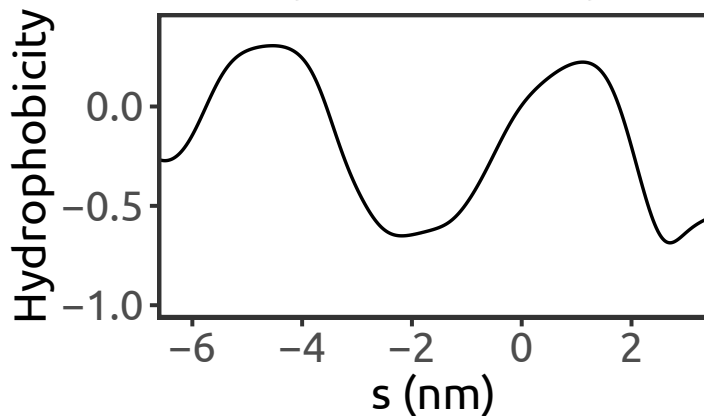
X-ray (3.5 Å)

Hassaine et al., 2014

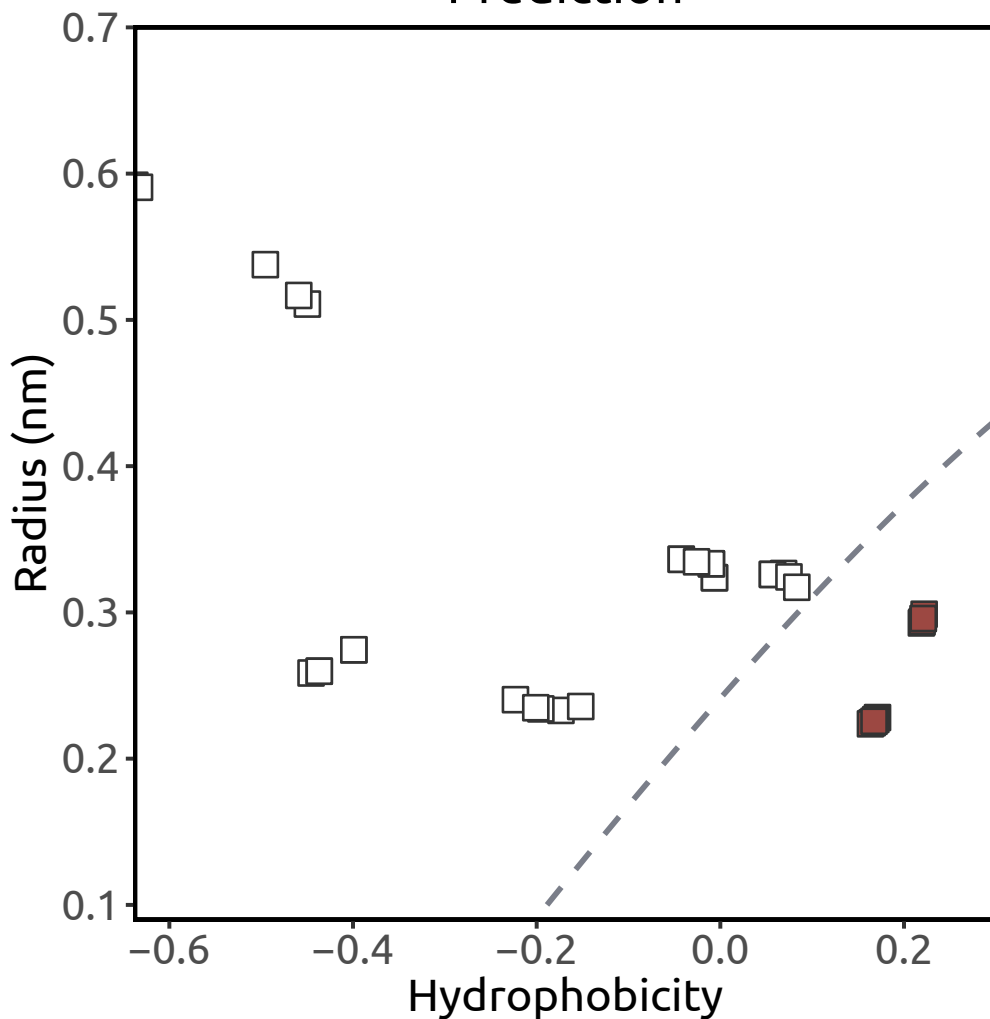
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.93 (n = 10)

Simulation result:
barrier to water

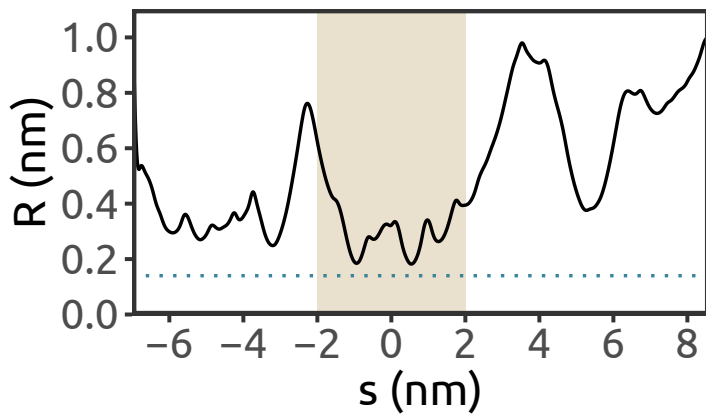
5HT3R (PDB ID: 6BE1)

Mus musculus

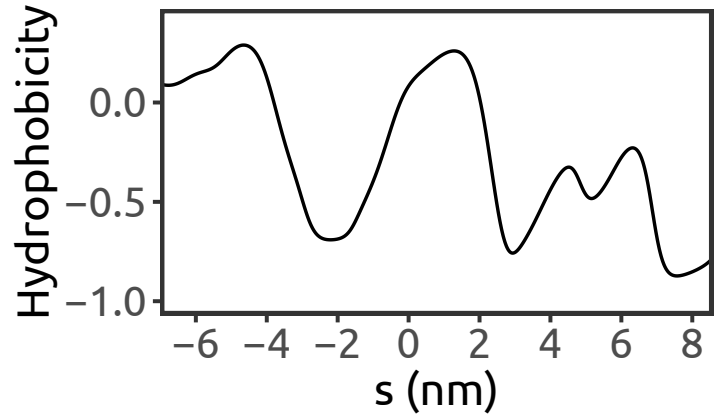
cryo-EM (4.31 Å)

Basak et al., 2018

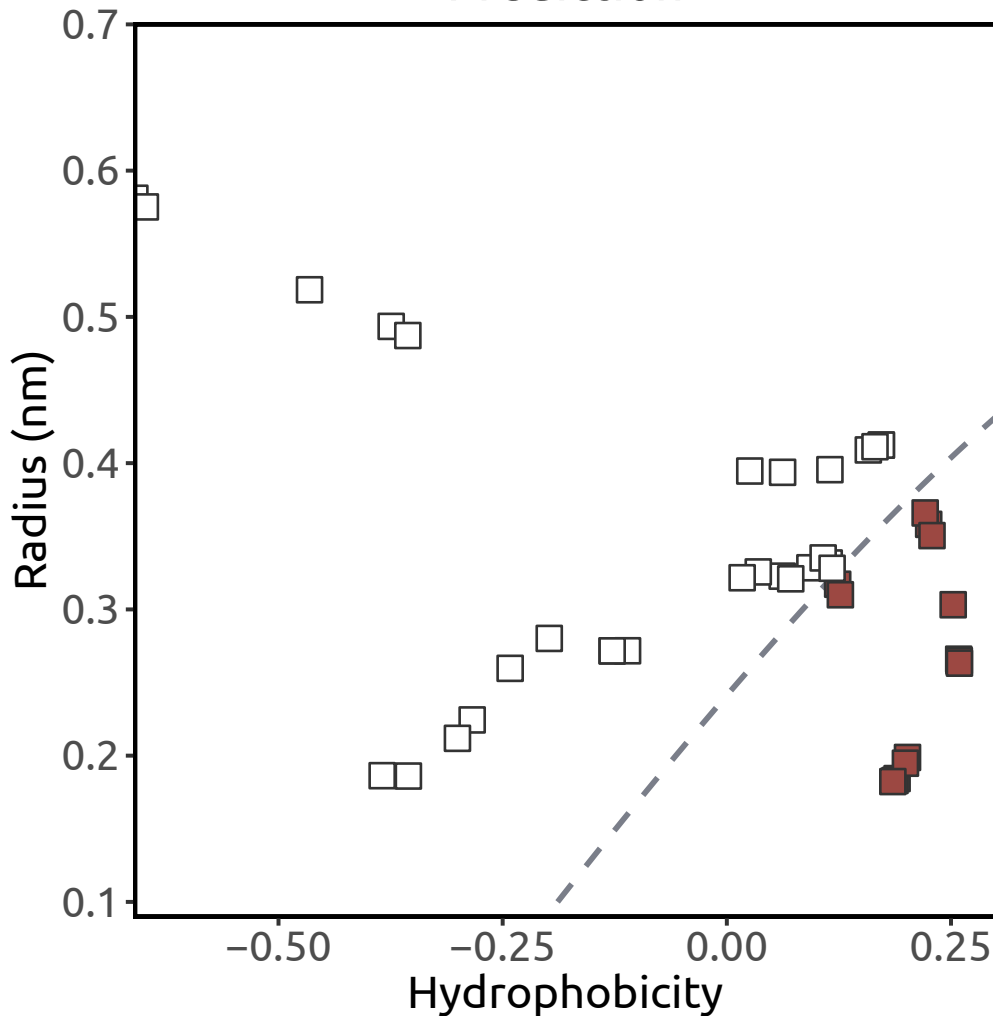
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.45 (n = 15)

Simulation result:
barrier to water

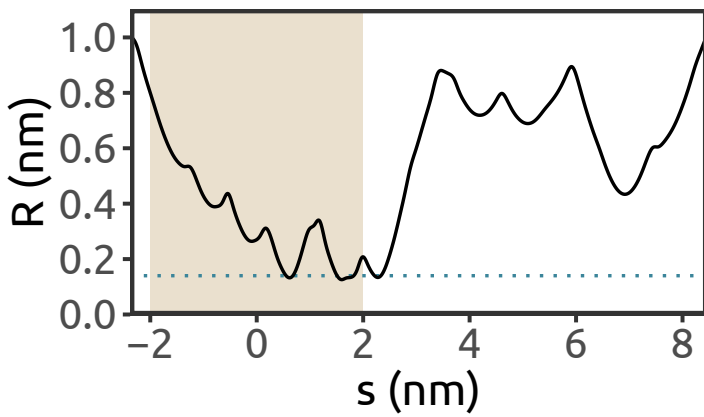
ELIC (PDB ID: 2VL0)

Erwinia chrysanthemi

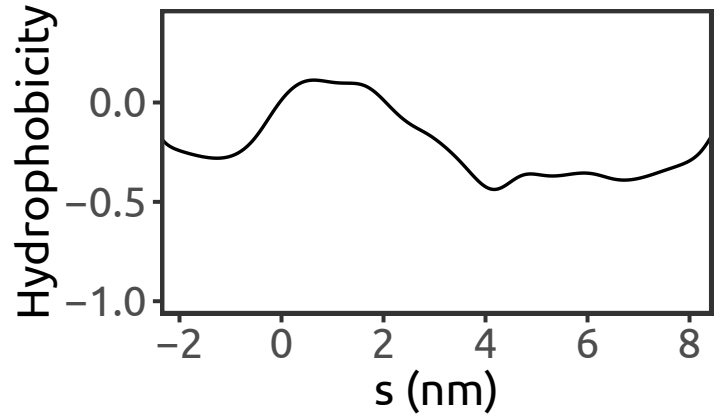
X-ray (3.3 Å)

Hilf & Dutzler, 2008

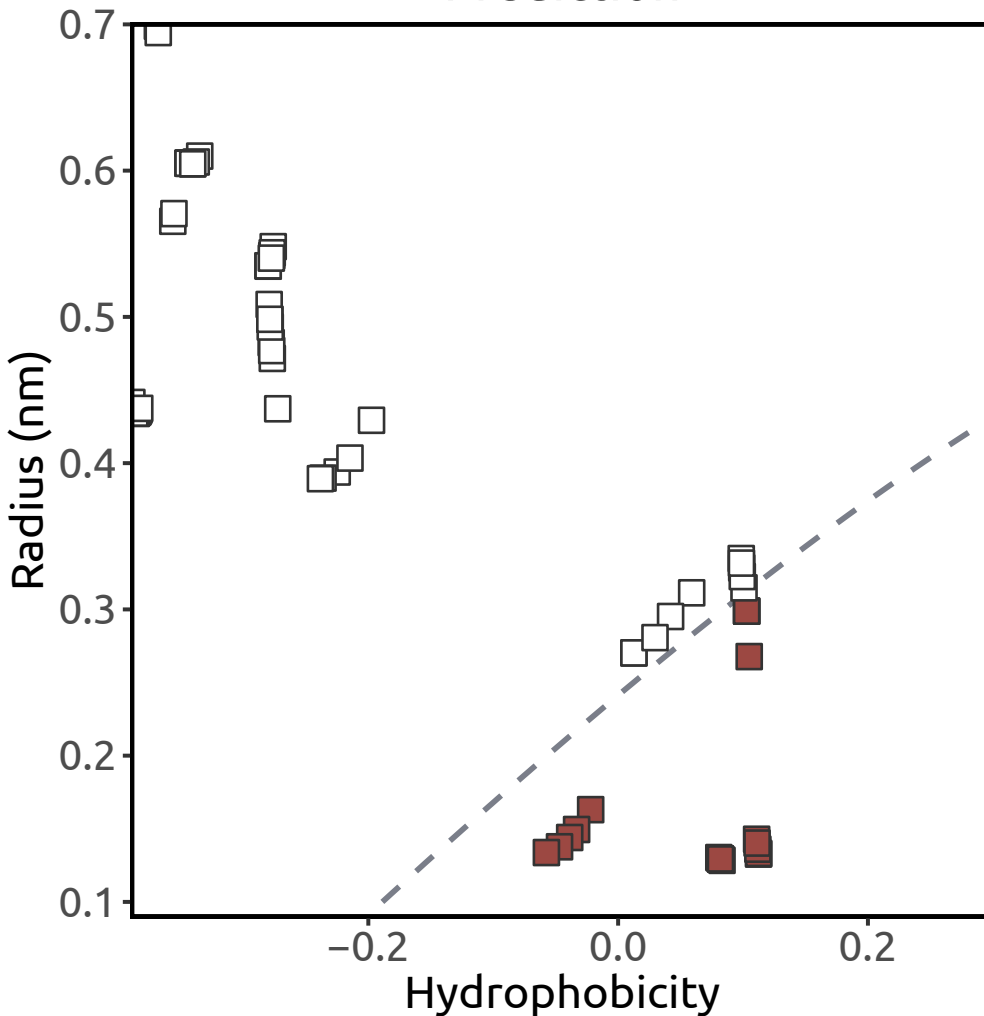
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.78 (n = 19)

Simulation result:

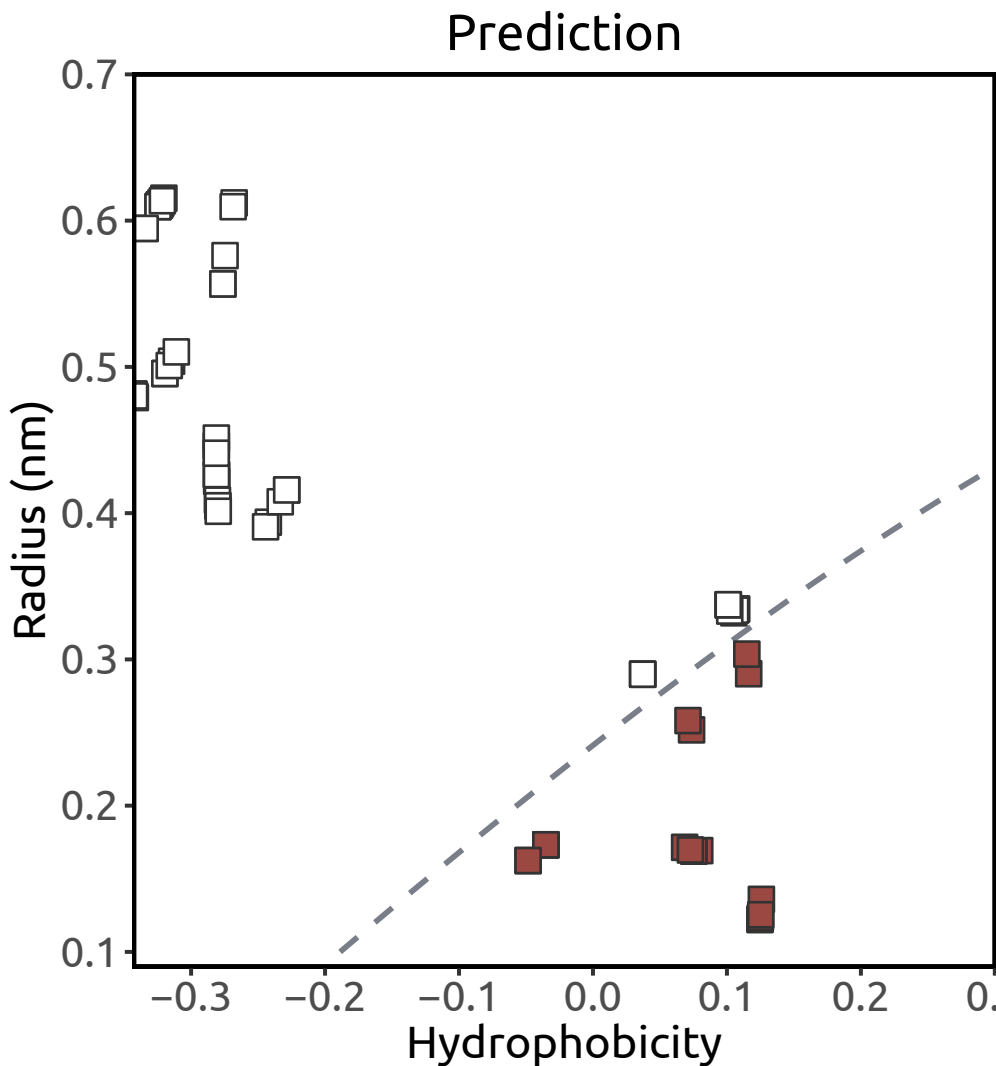
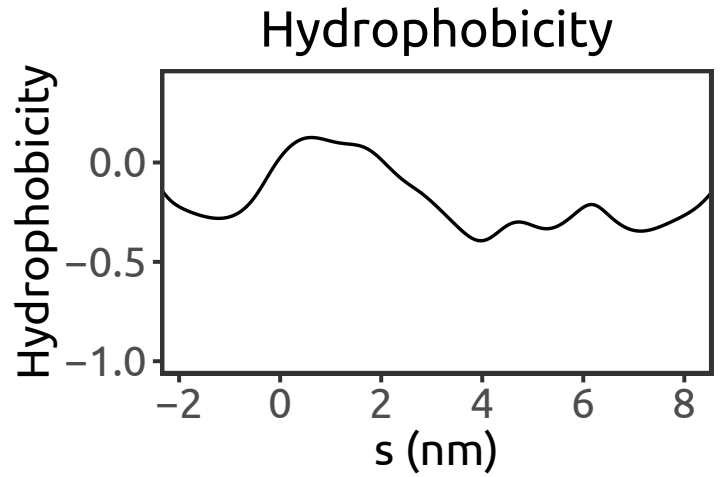
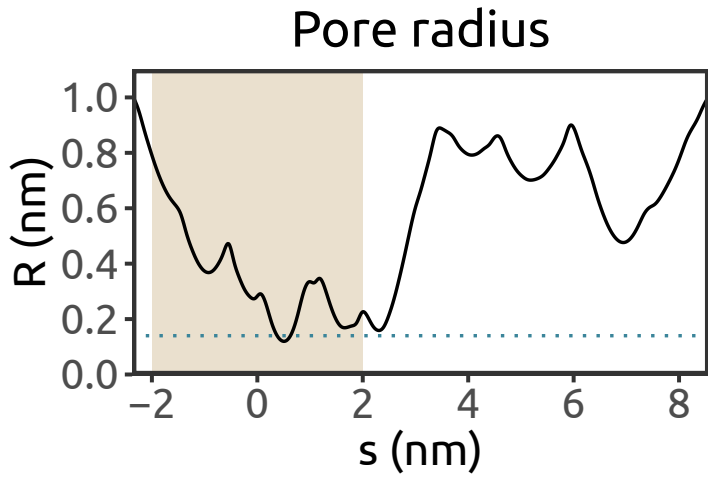
barrier to water

ELIC (PDB ID: 3RQW)

Erwinia chrysanthemi

X-ray (2.91 Å)

Pan et al., 2012



Heuristic score:

1.61 (n = 19)

Simulation result:

barrier to water

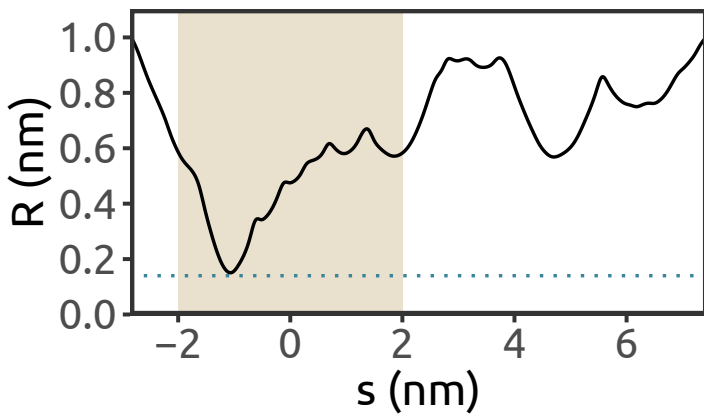
GABAAR (PDB ID: 4COF)

Homo sapiens

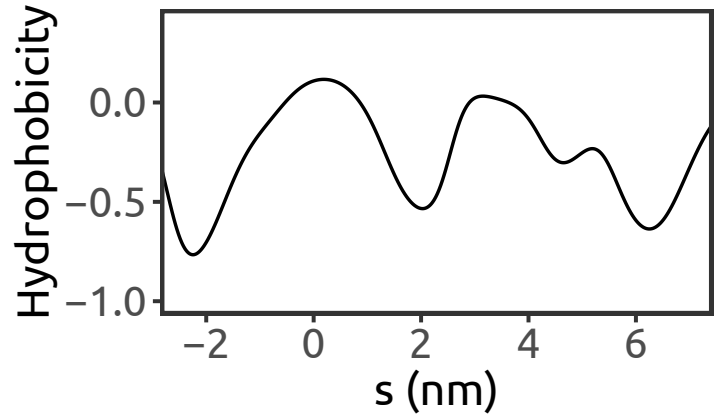
X-ray (2.97 Å)

Miller et al., 2014

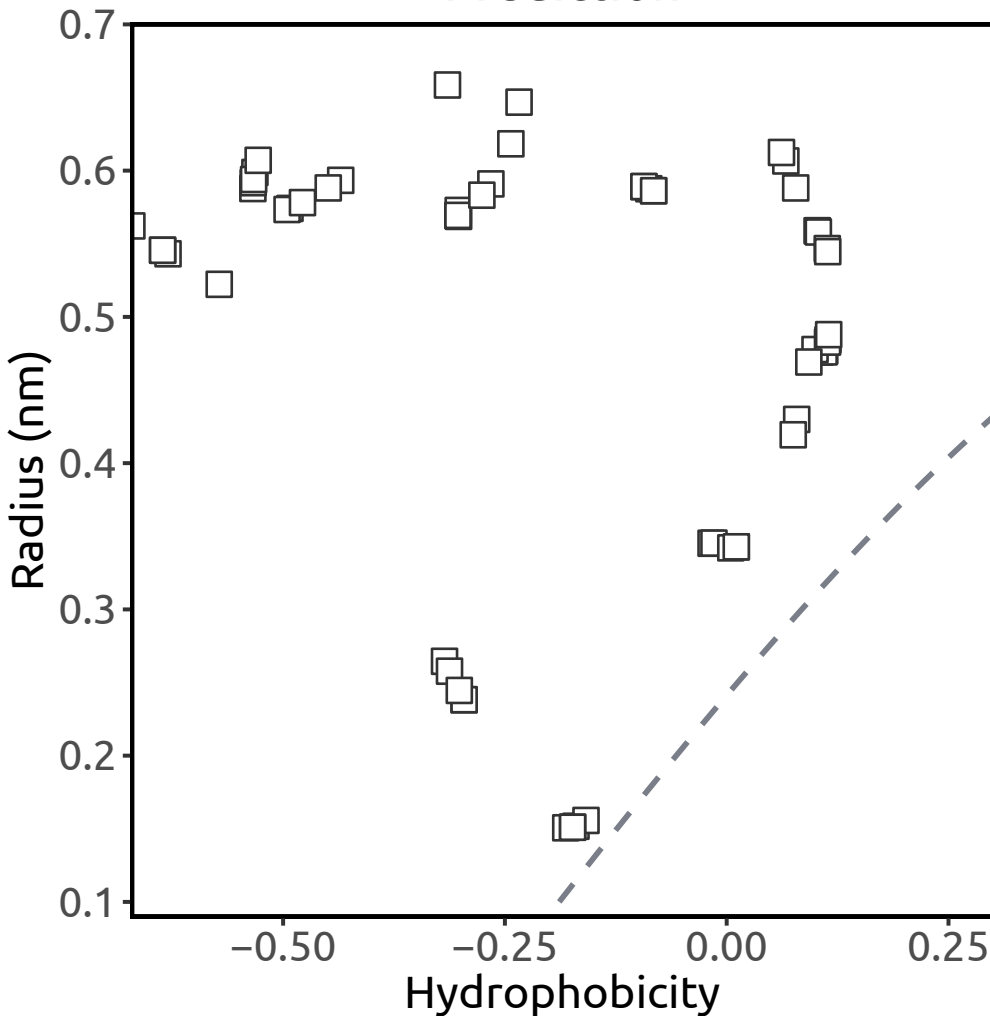
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0 (n = 0)

Simulation result:

hydrated channel

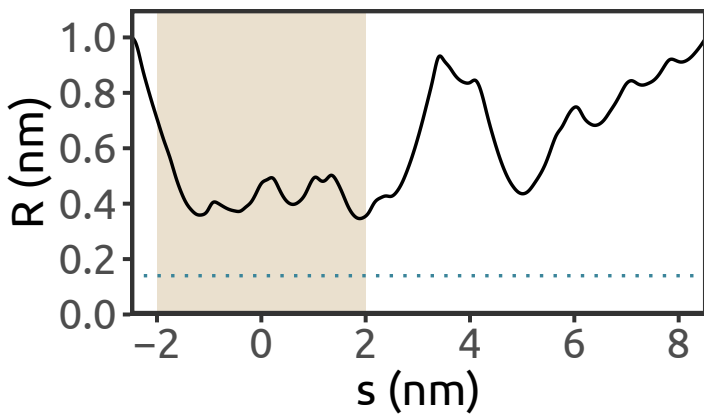
GABAAR (PDB ID: 6A96)

Homo sapiens

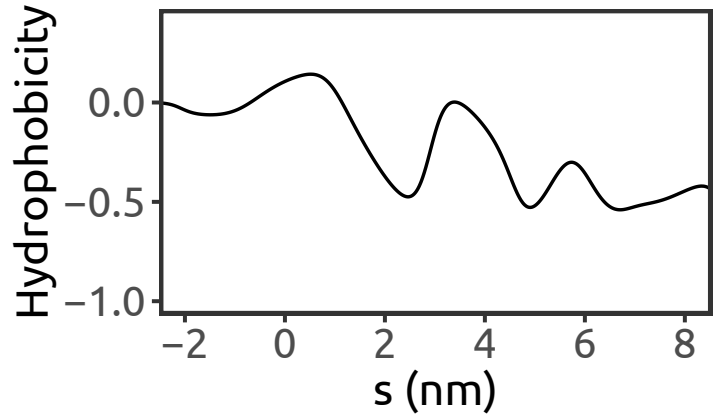
cryo-EM (3.51 Å)

Liu et al., 2018

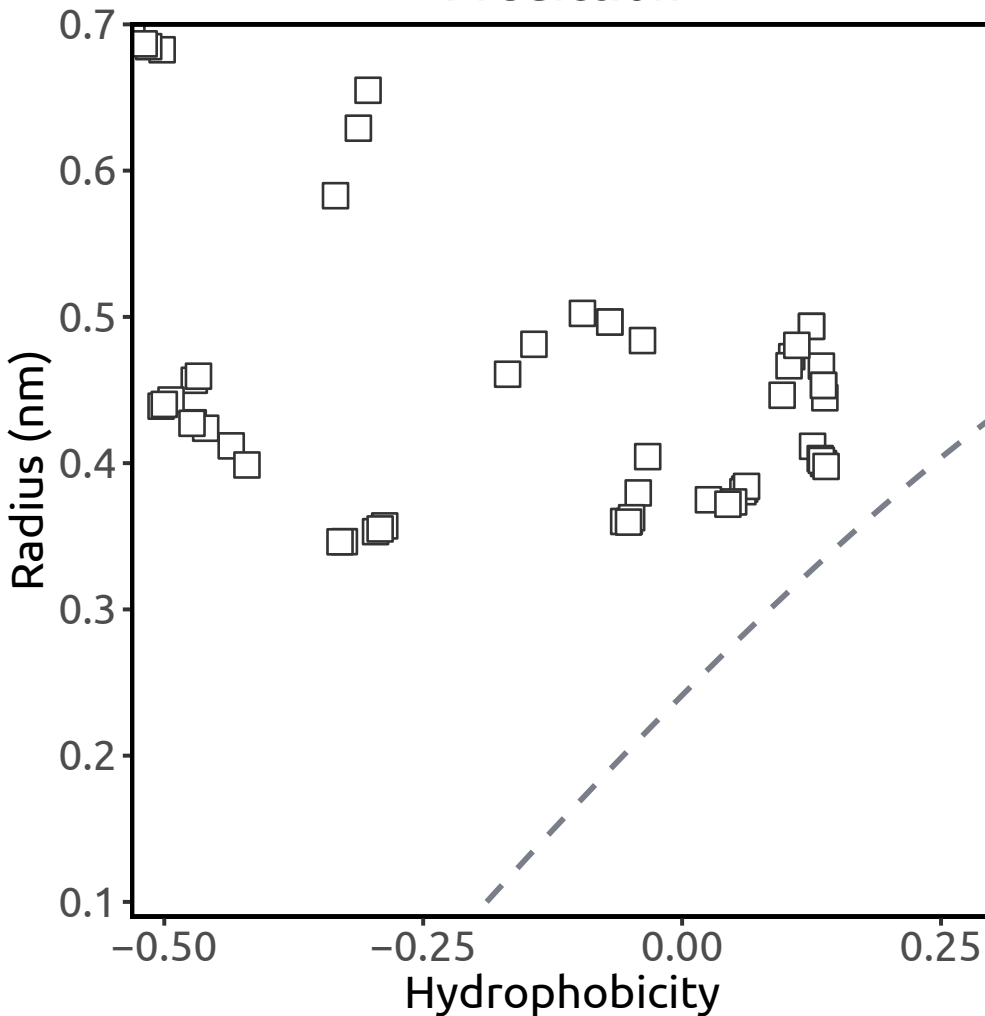
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0 (n = 0)

Simulation result:

hydrated channel

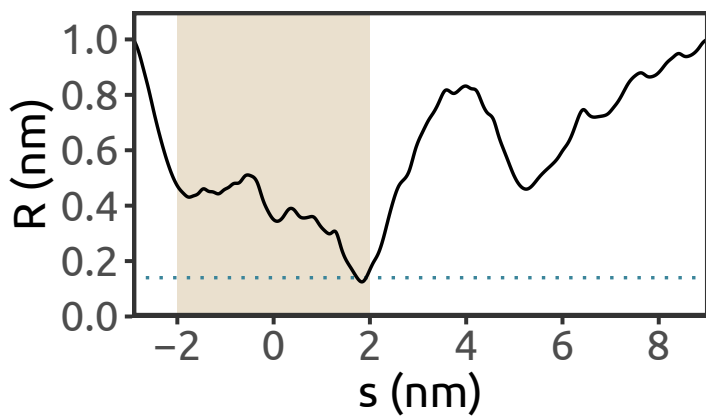
GABAAR (PDB ID: 6D6T)

Homo sapiens

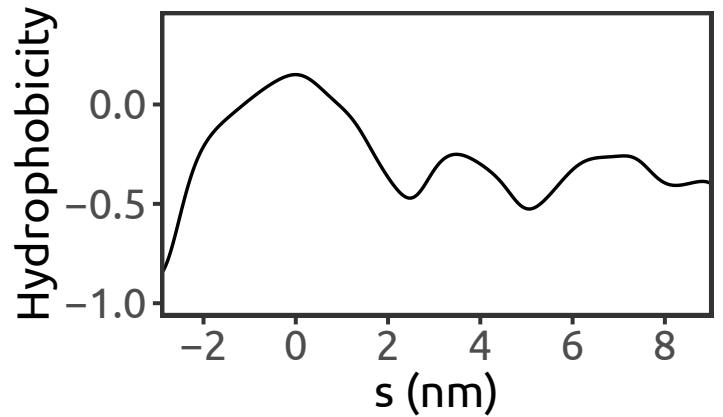
cryo-EM (3.86 Å)

Zhu et al., 2018

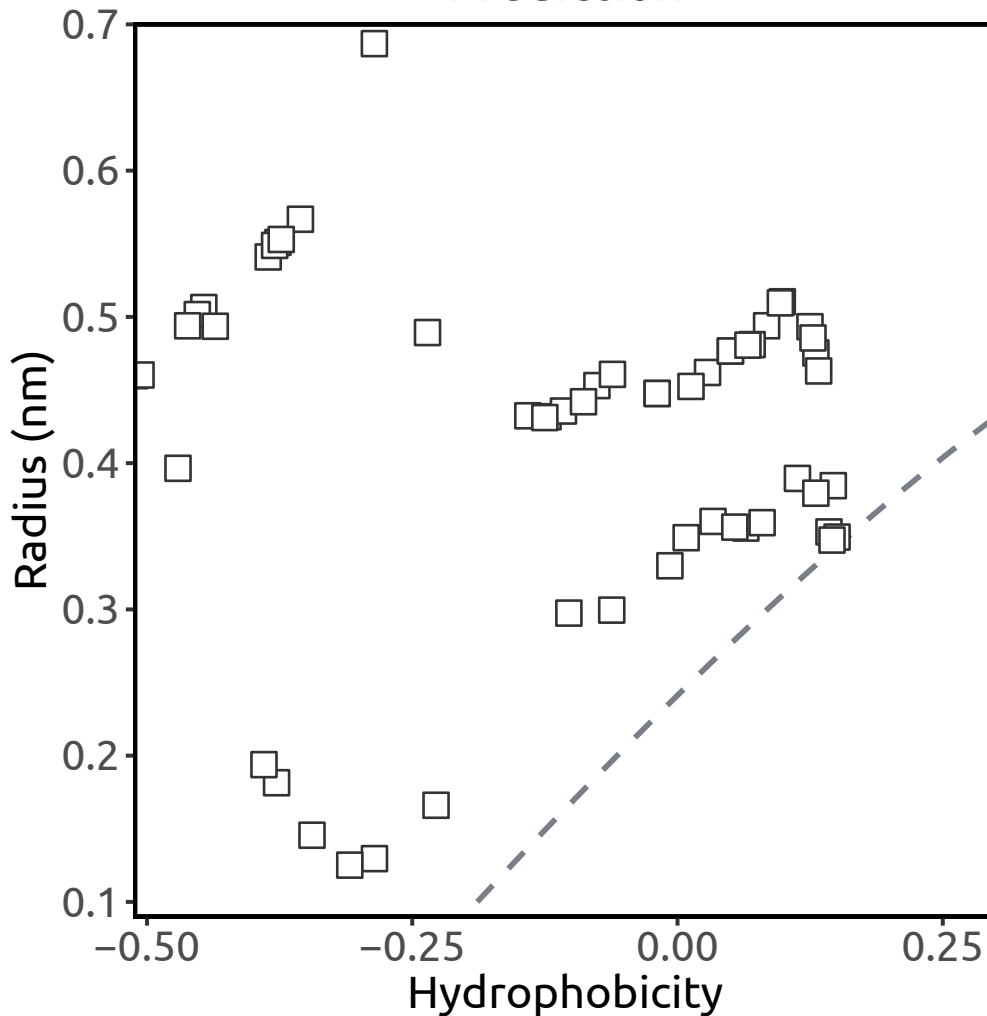
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 ($n = 0$)

Simulation result:
hydrated channel

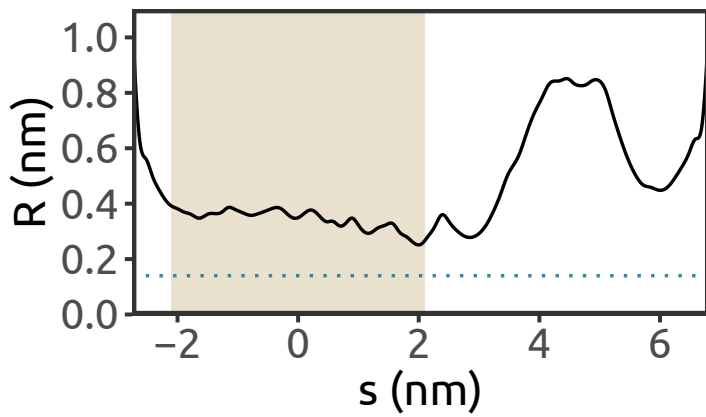
GABAAR (PDB ID: 6D6U)

Homo sapiens

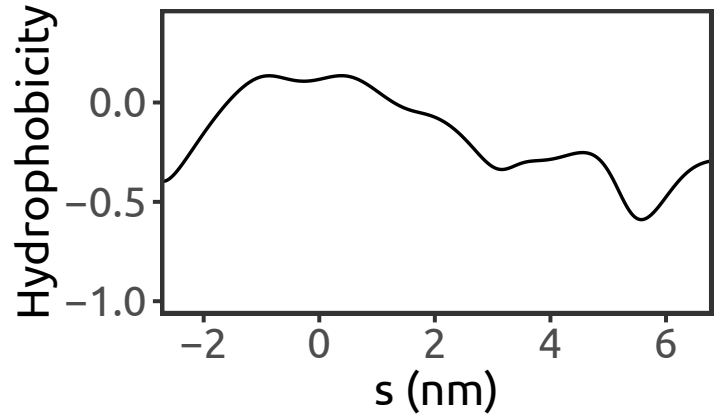
cryo-EM (3.92 Å)

Zhu et al., 2018

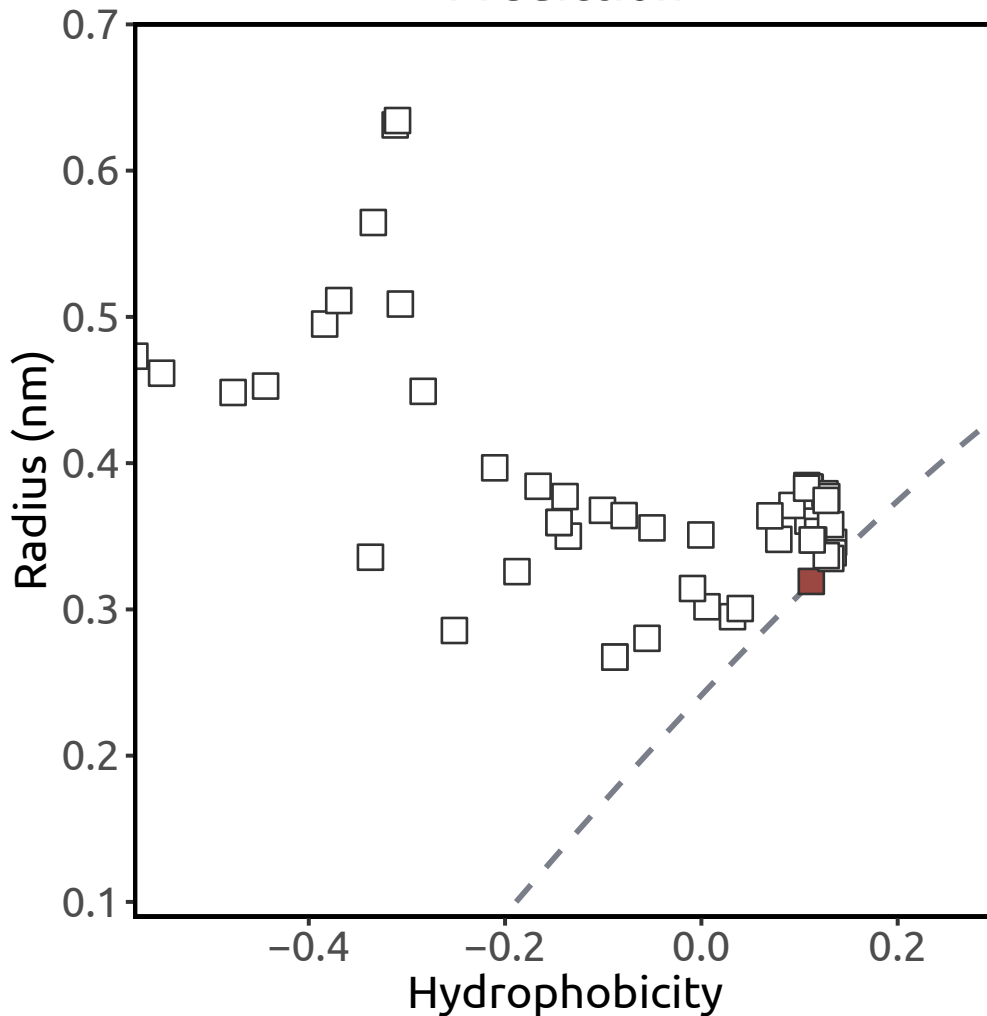
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 1)

Simulation result:
barrier to water

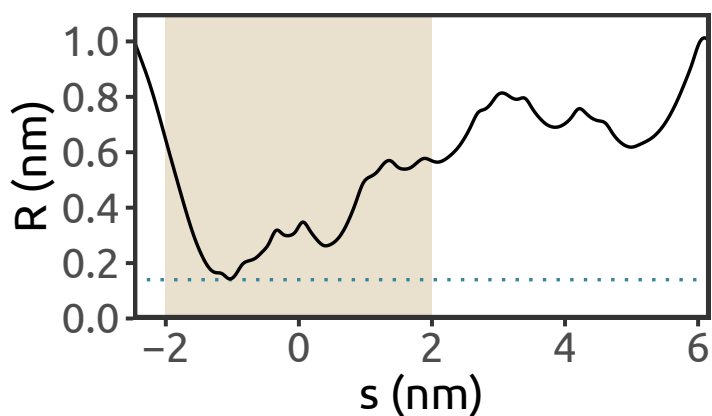
GLIC (PDB ID: 2XQ7)

Gloeobacter violaceus

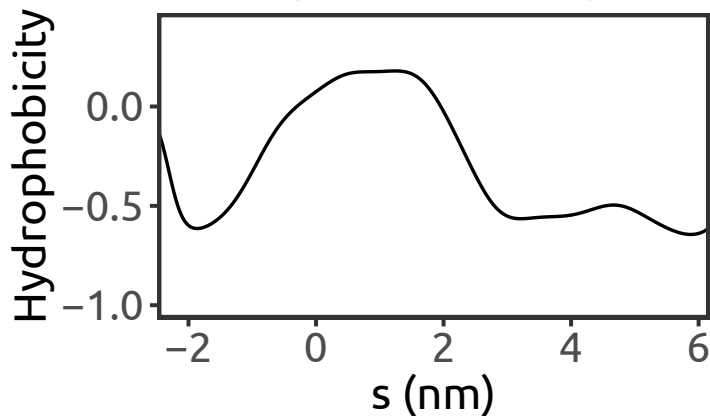
X-ray (3.4 Å)

Hilf et al., 2010

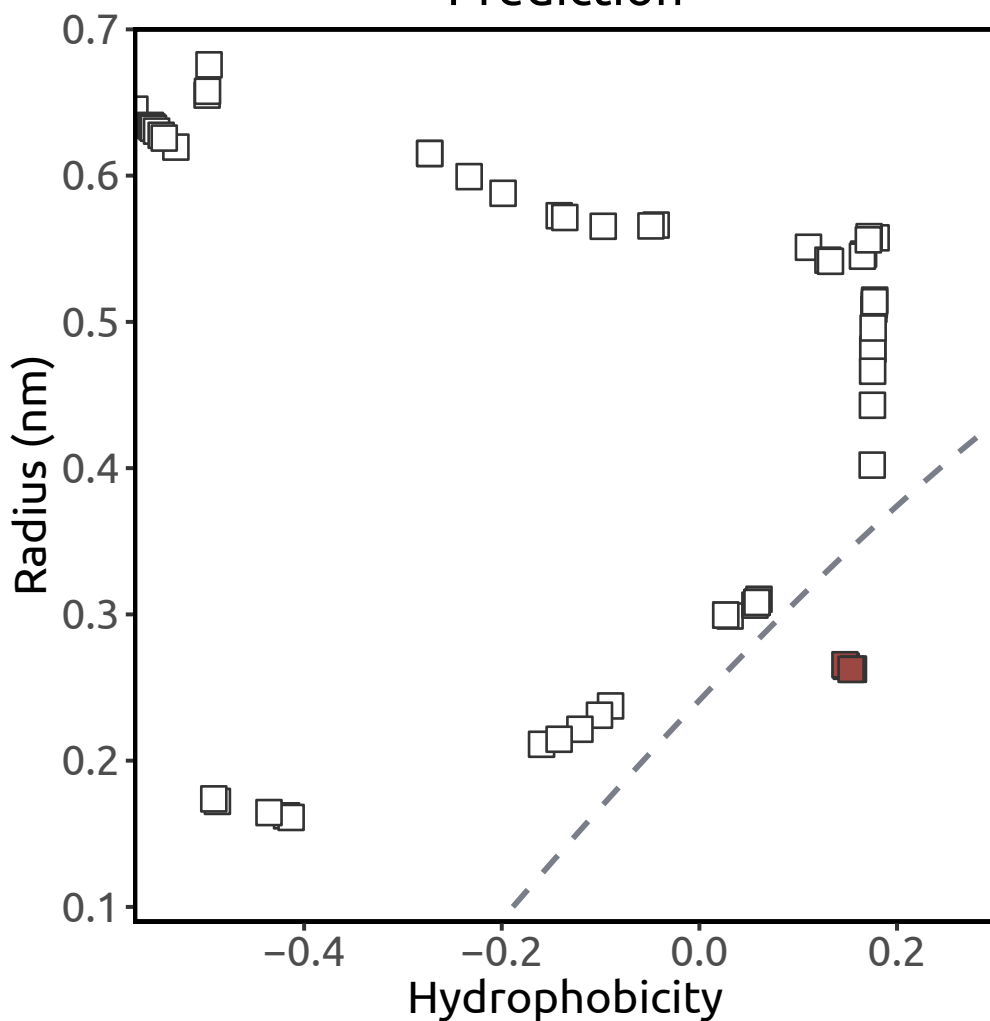
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.34 (n = 5)

Simulation result:
hydrated channel

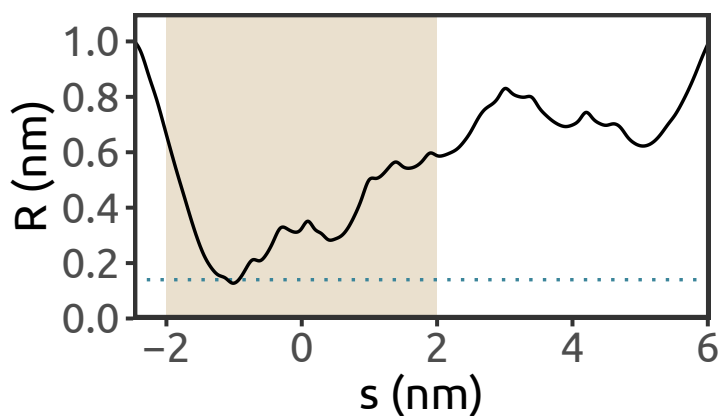
GLIC (PDB ID: 3EHZ)

Gloeobacter violaceus

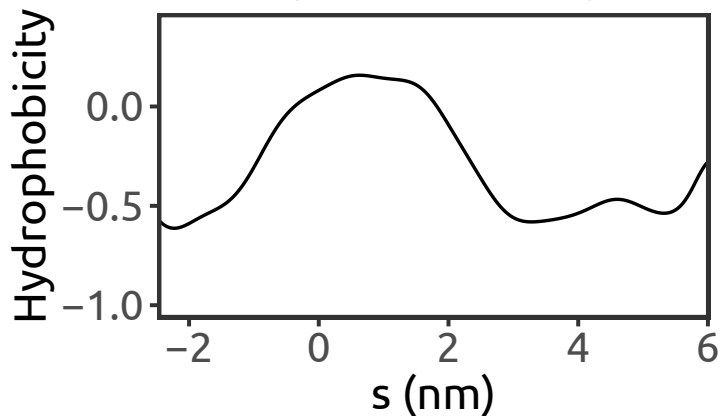
X-ray (3.1 Å)

Hilf & Dutzler, 2009

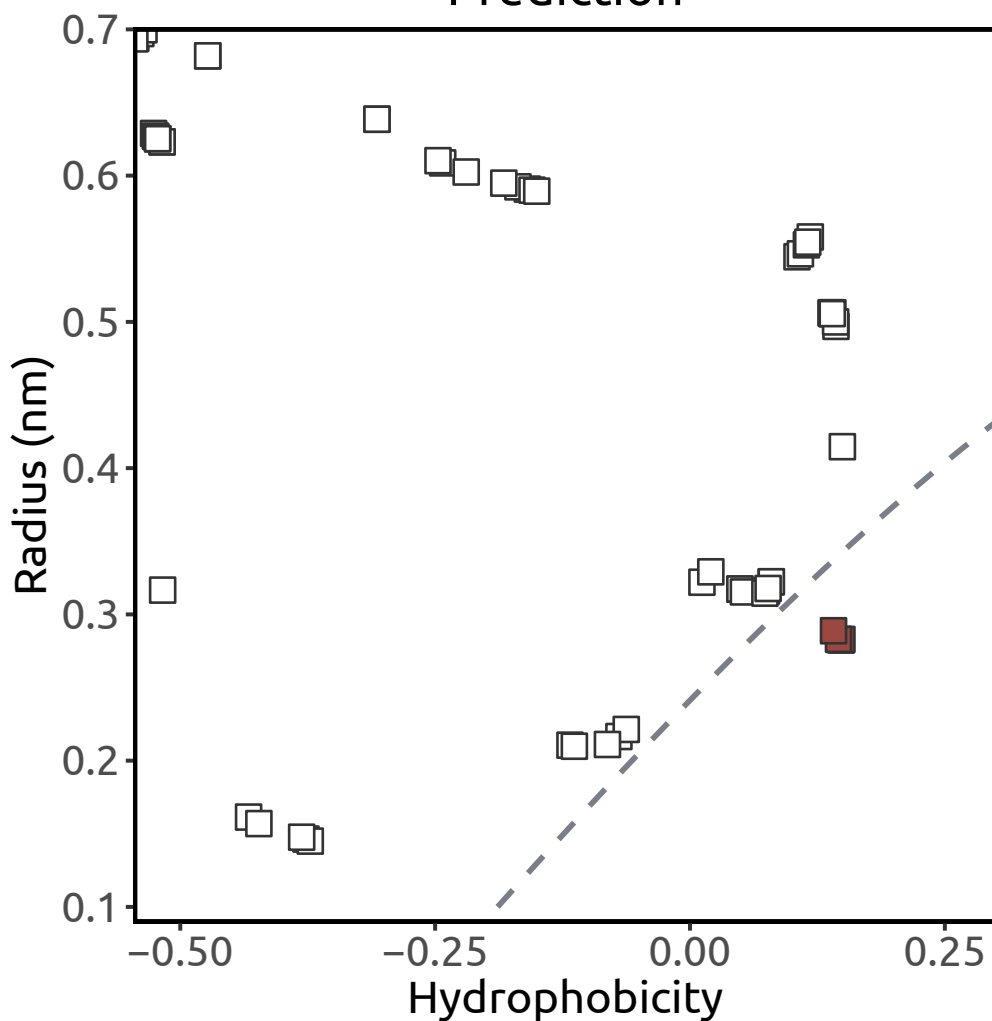
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.24 ($n = 5$)

Simulation result:
hydrated channel

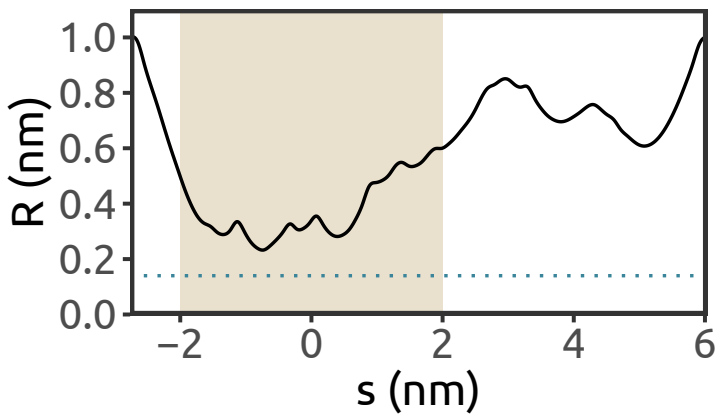
GLIC (PDB ID: 4F8H)

Gloeobacter violaceus

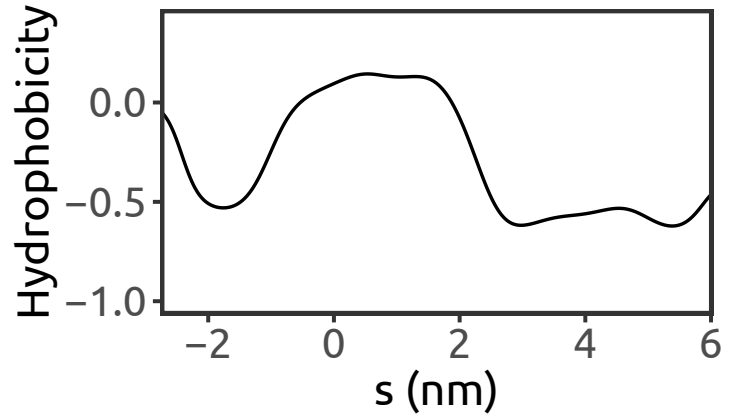
X-ray (2.99 Å)

Pan et al., 2012

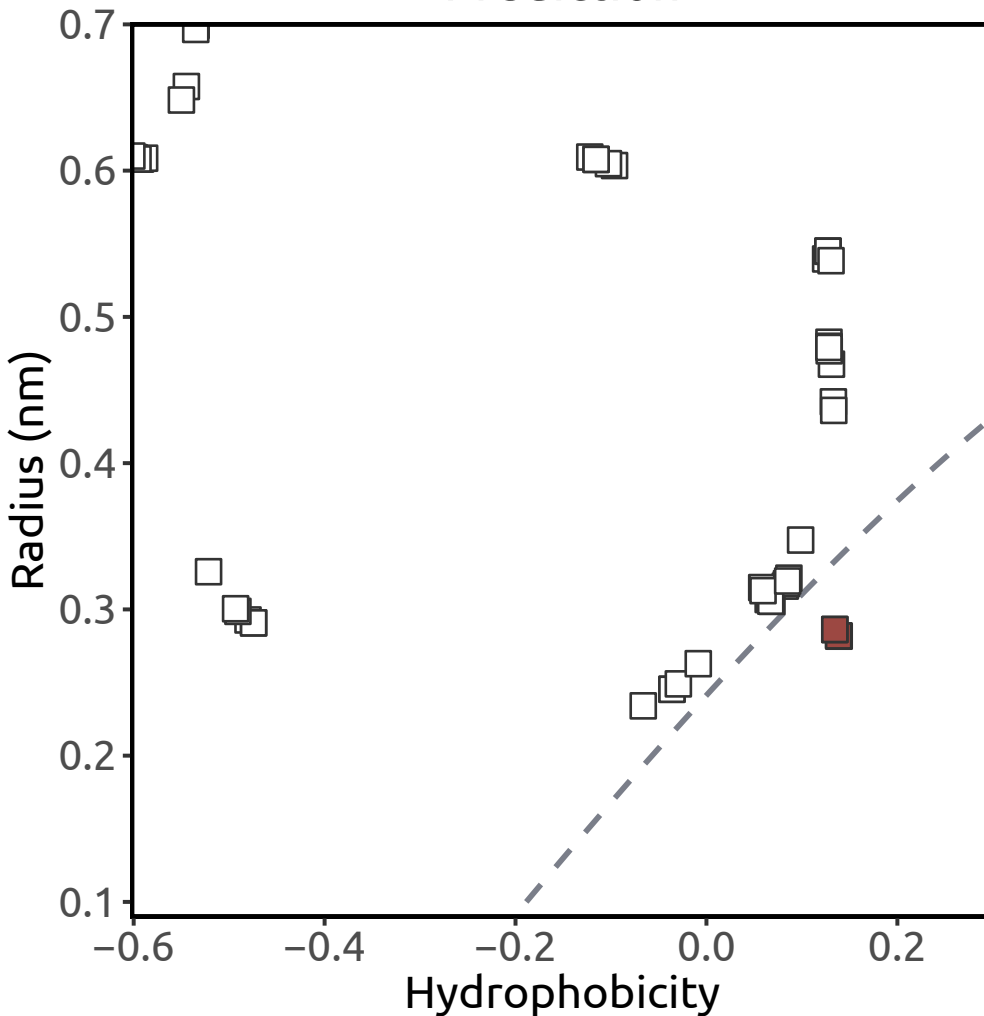
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.21 (n = 5)

Simulation result:
hydrated channel

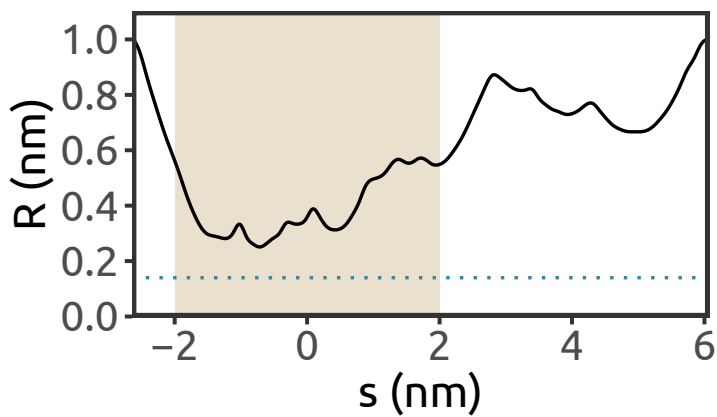
GLIC (PDB ID: 4HFI)

Gloeobacter violaceus

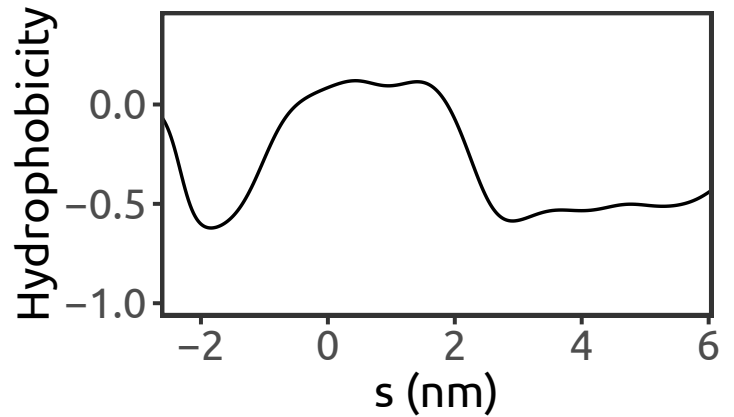
X-ray (2.4 Å)

Sanguet et al., 2013

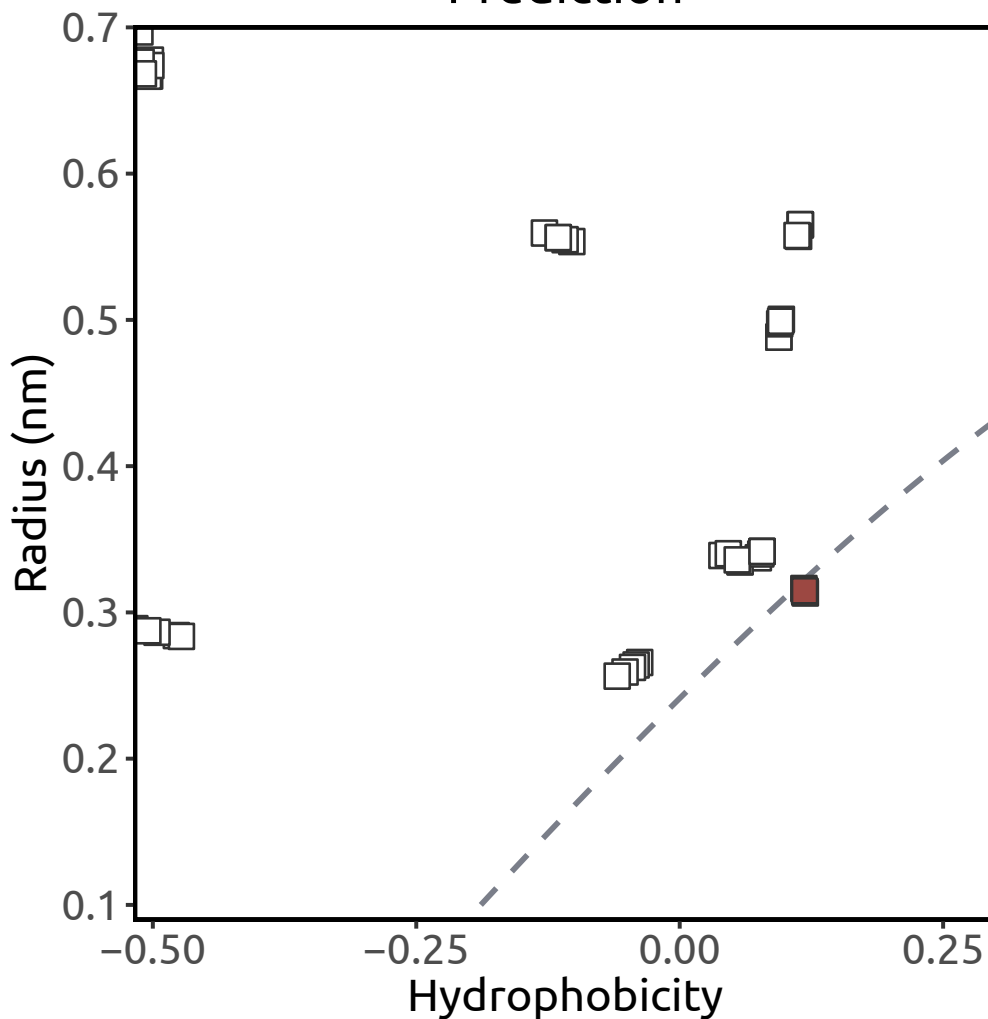
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.04 ($n = 5$)

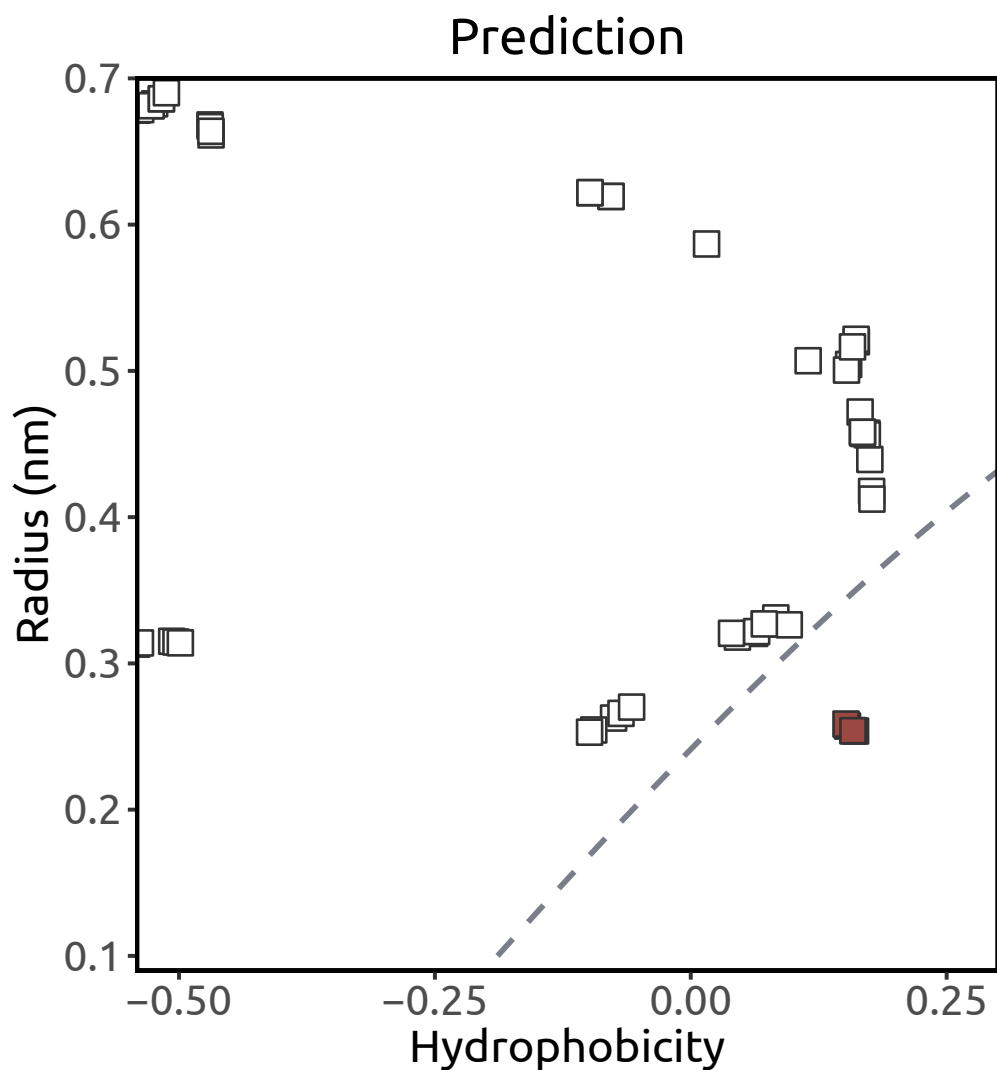
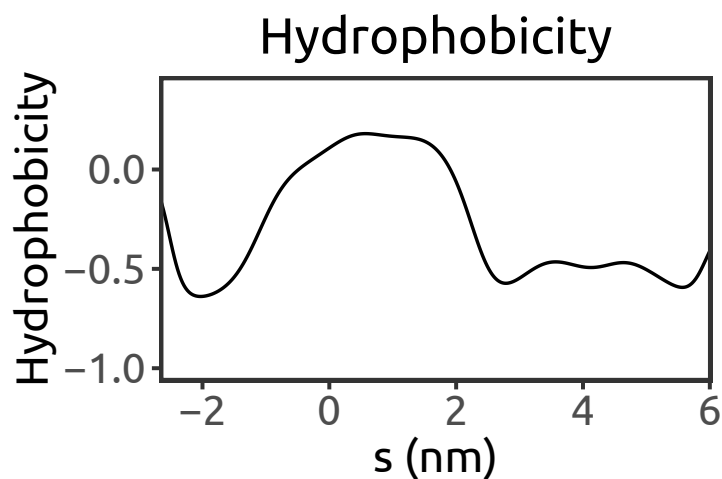
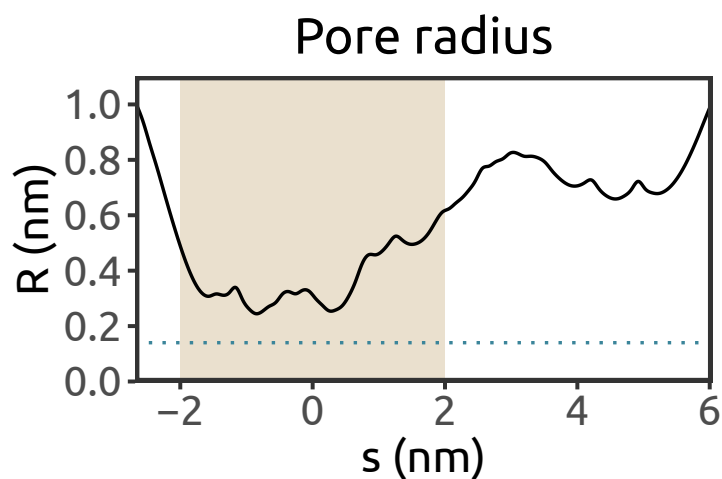
Simulation result:
hydrated channel

GLIC (PDB ID: 4NPP)

Gloeobacter violaceus

X-ray (3.35 Å)

Sanguet et al., 2014



Heuristic score:
0.39 (n = 5)

Simulation result:
hydrated channel

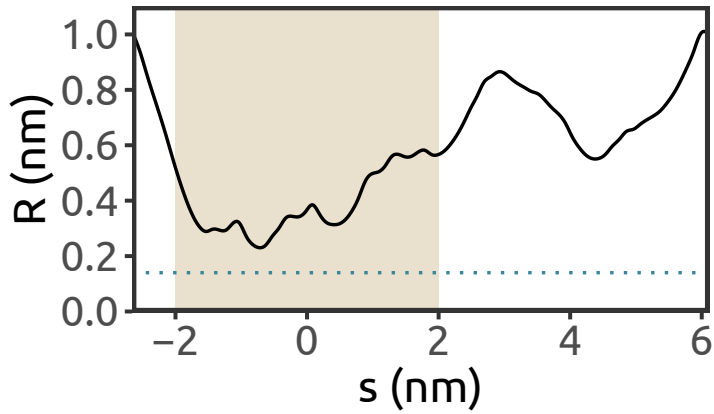
GLIC (PDB ID: 4QH5)

Gloeobacter violaceus

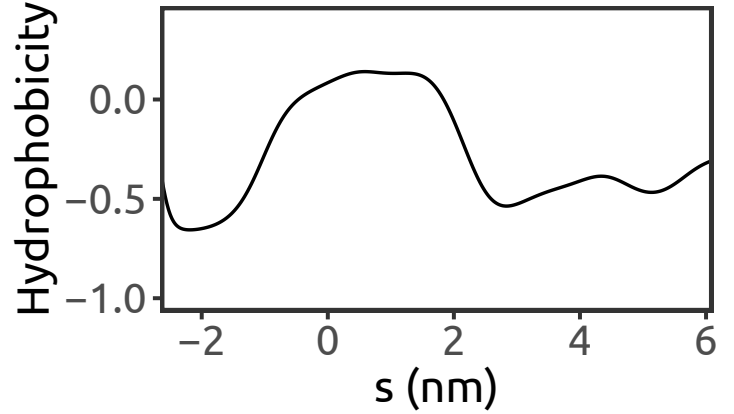
X-ray (3 Å)

Fourati et al., 2015

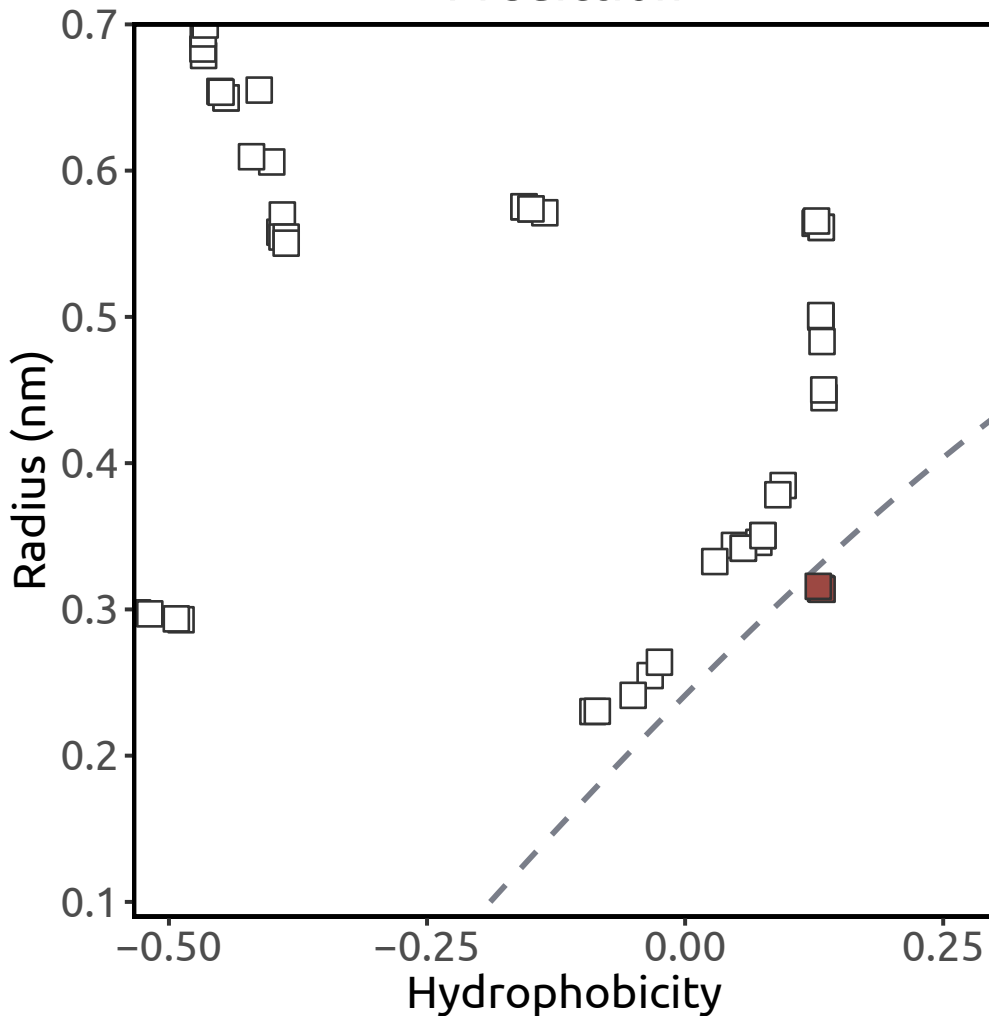
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.07 ($n = 5$)

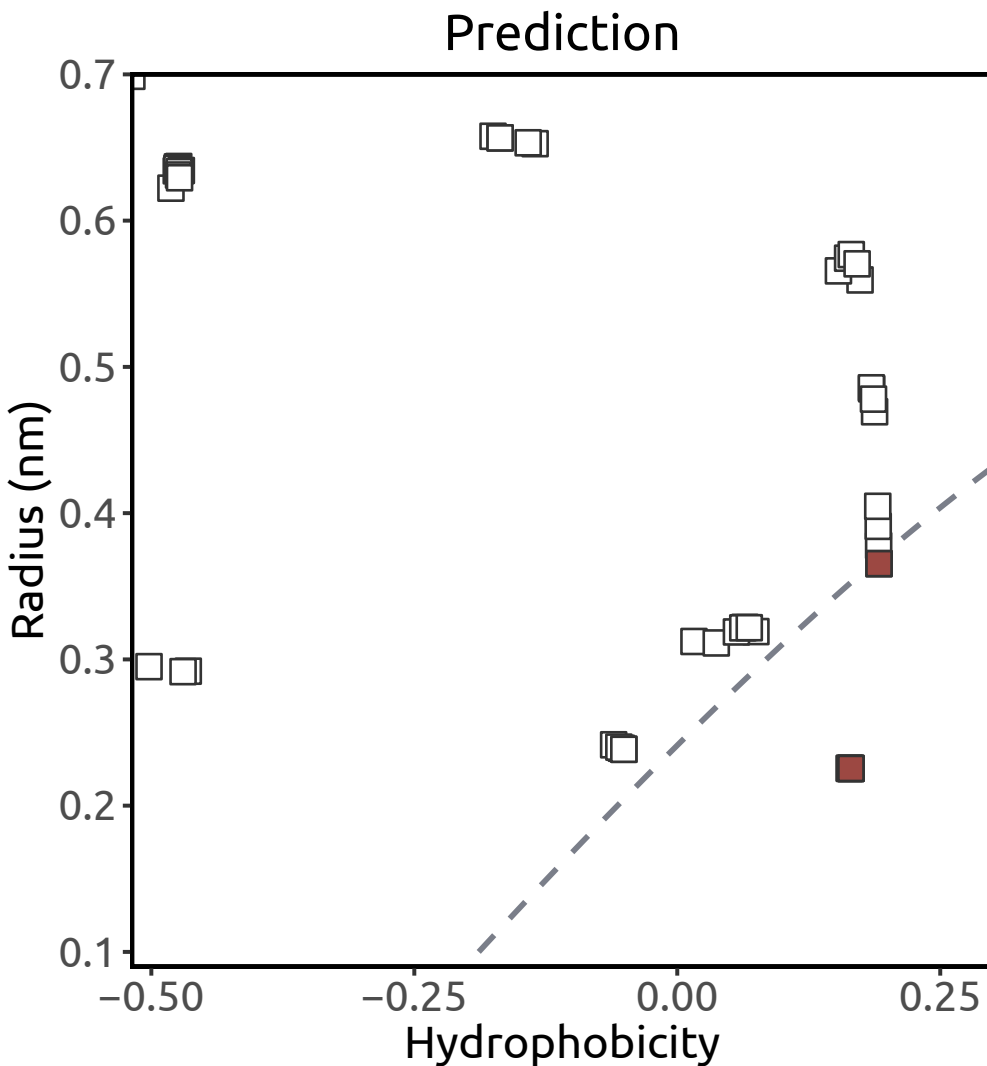
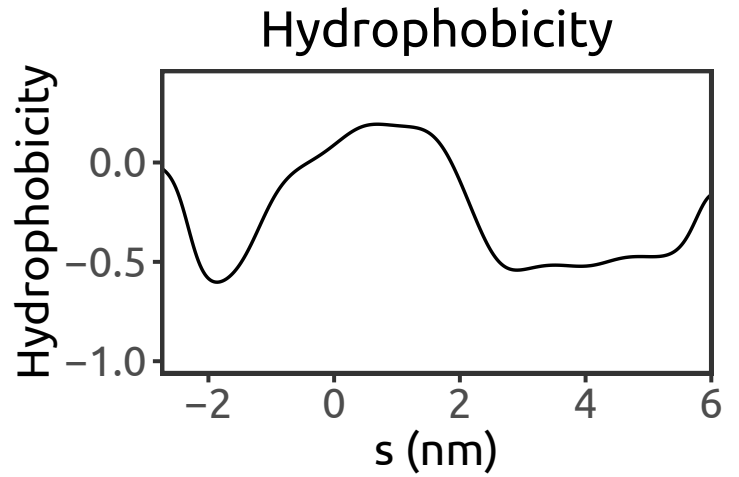
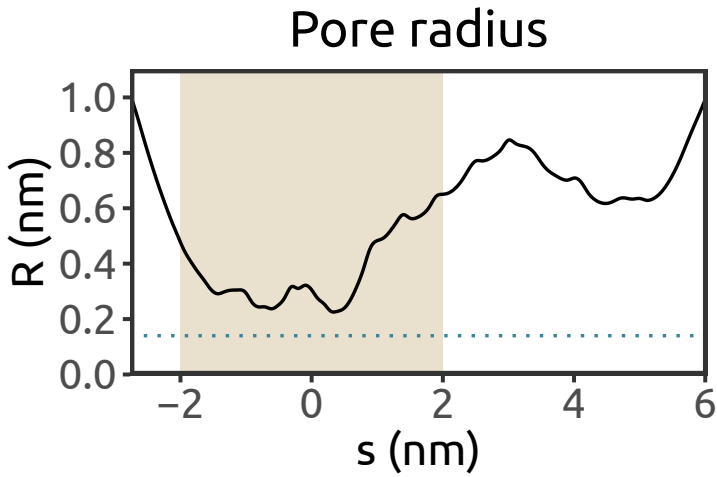
Simulation result:
hydrated channel

GLIC (PDB ID: 5J0Z)

Gloeobacter violaceus

X-ray (3.25 Å)

Basak et al., 2017



Heuristic score:
0.54 ($n = 7$)

Simulation result:
hydrated channel

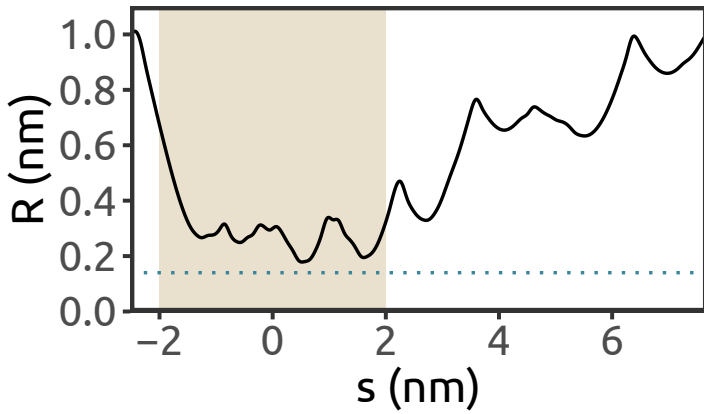
GLIC (PDB ID: 5L47)

Gloeobacter violaceus

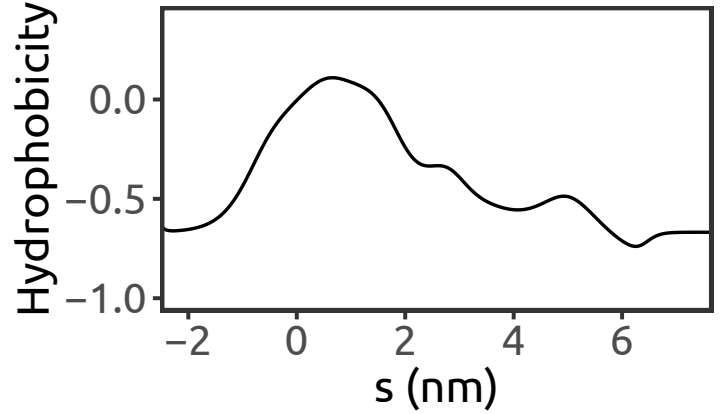
X-ray (3.3 Å)

Fourati et al., 2017

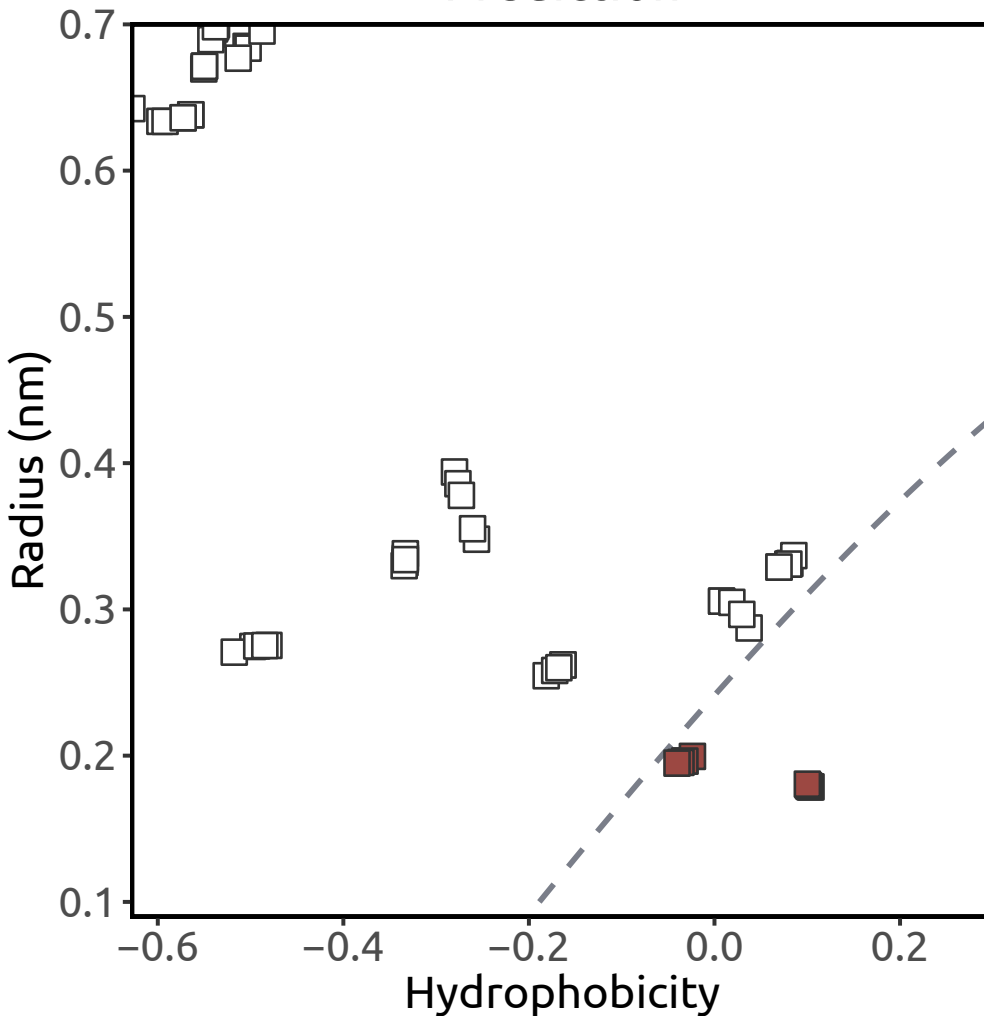
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.63 (n = 10)

Simulation result:

barrier to water

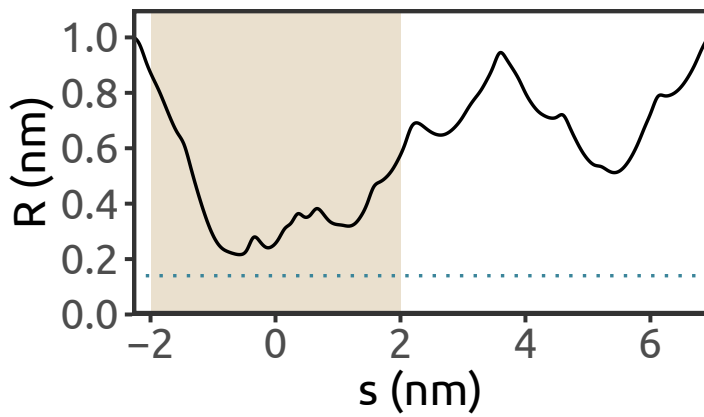
GluCl (PDB ID: 3RHW)

Caenorhabditis elegans

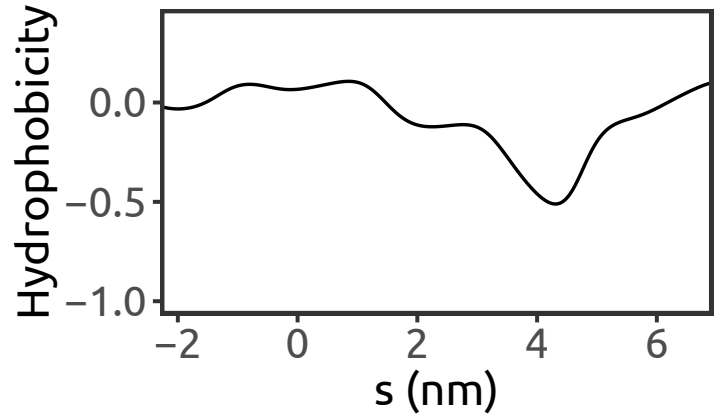
X-ray (3.26 Å)

Pan et al., 2012

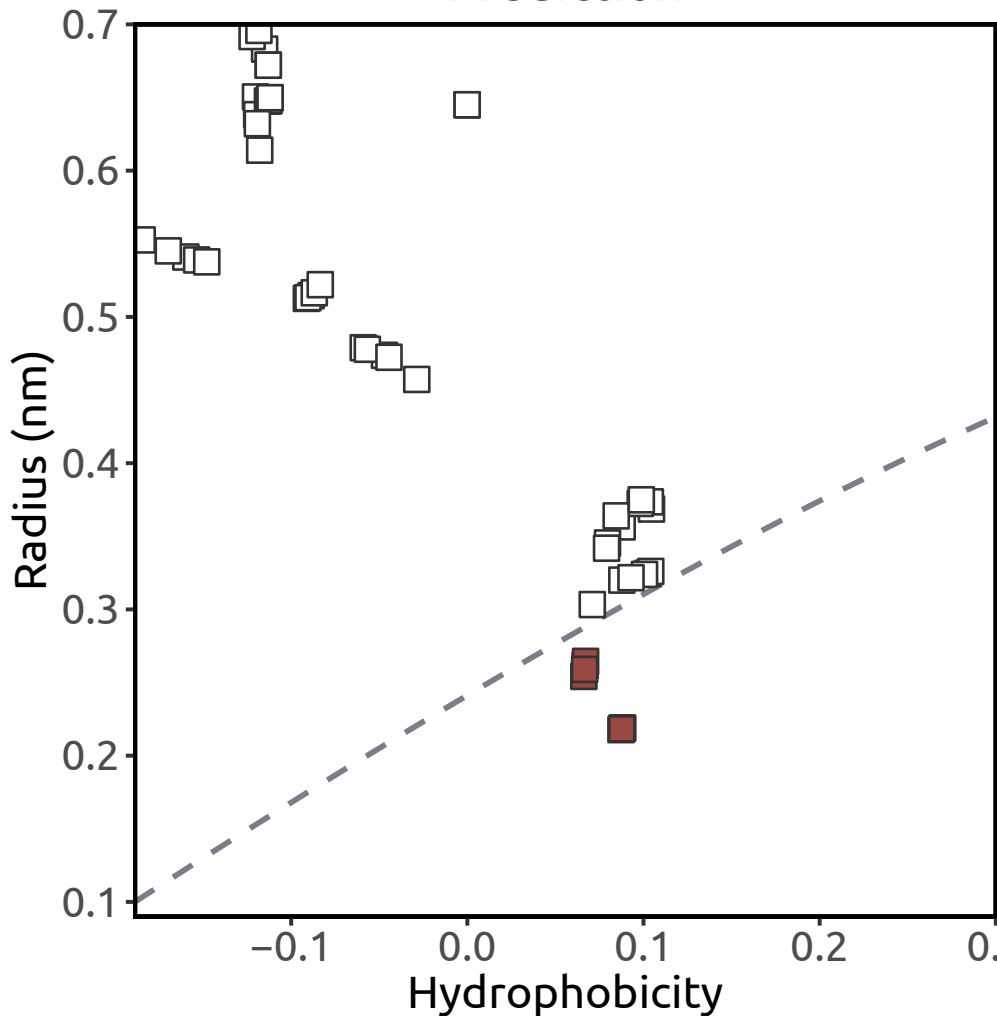
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.44 (n = 9)

Simulation result:
hydrated channel

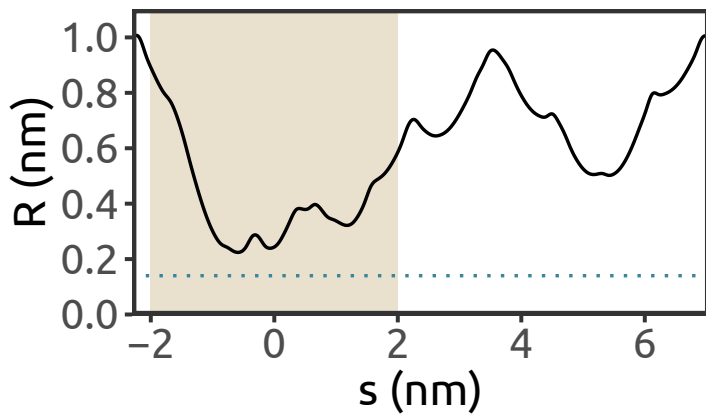
GluCl (PDB ID: 3RIF)

Caenorhabditis elegans

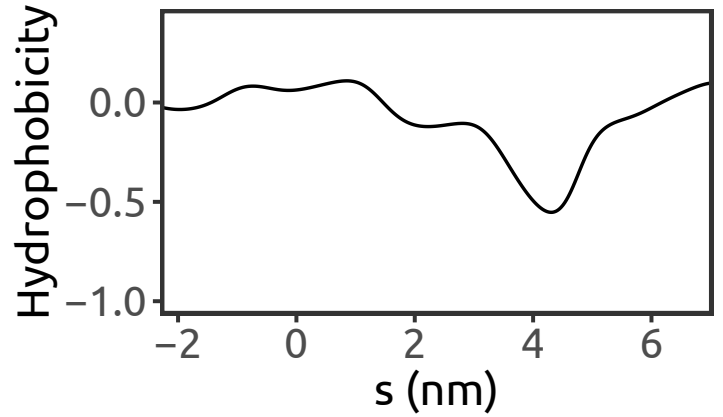
X-ray (3.35 Å)

Hibbs & Gounaux, 2011

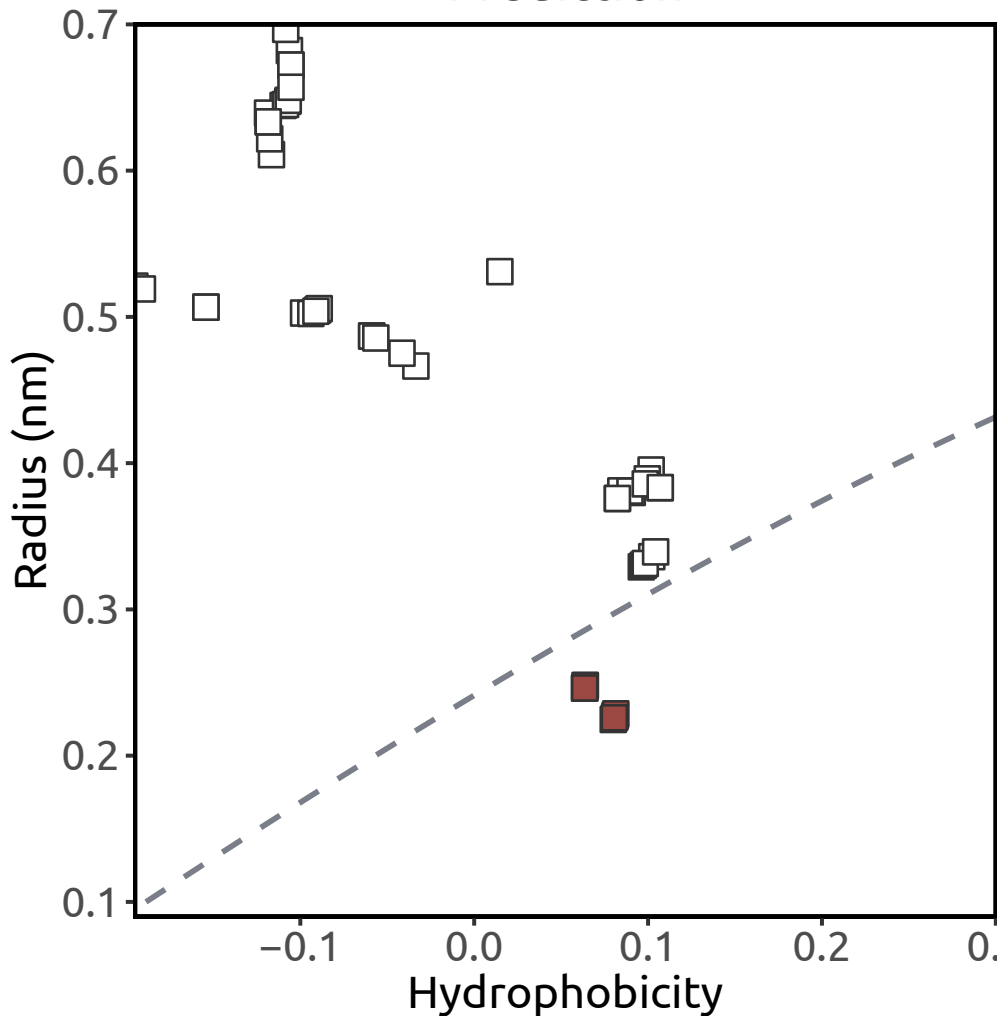
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.45 (n = 10)

Simulation result:
hydrated channel

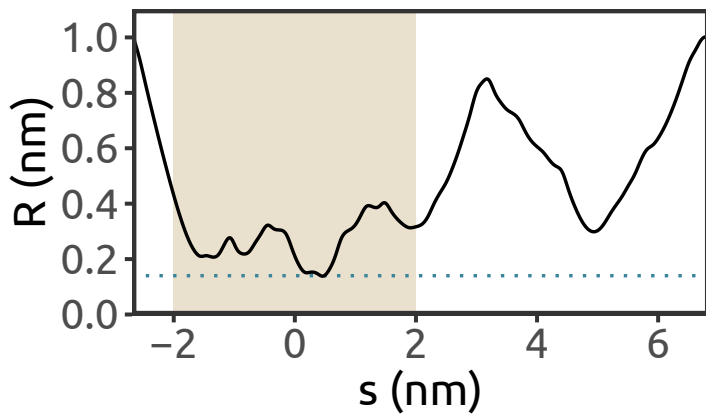
GluCl (PDB ID: 4TNV)

Caenorhabditis elegans

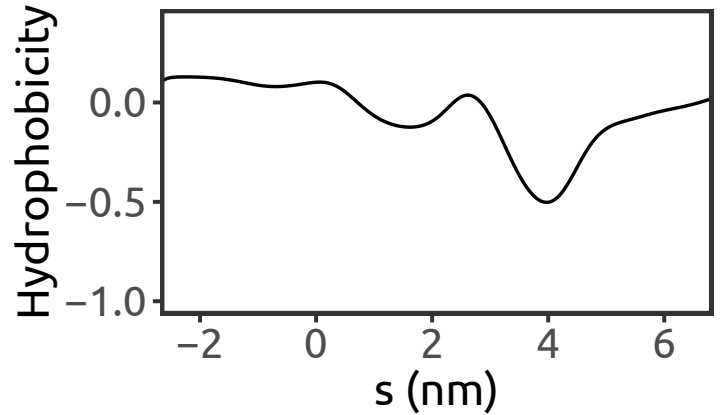
X-ray (3.6 Å)

Althoff et al., 2014

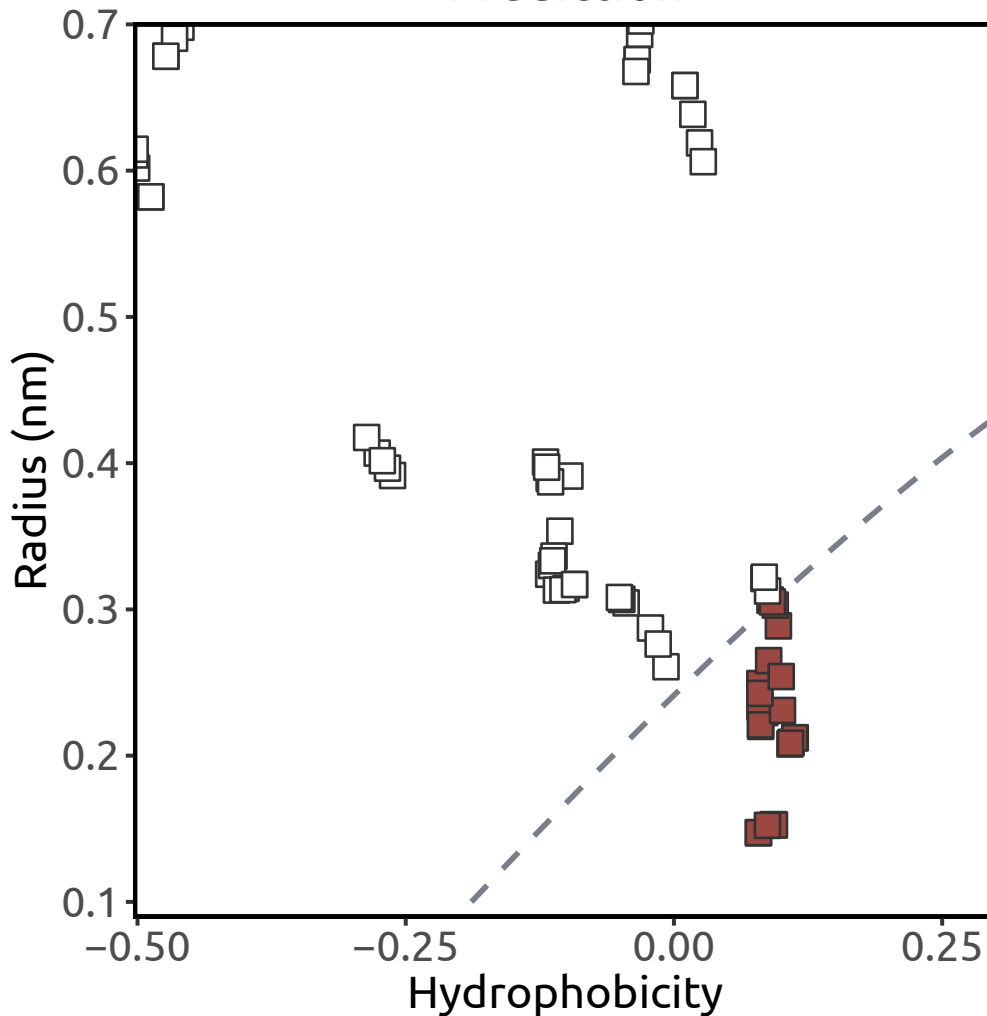
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.55 (n = 22)

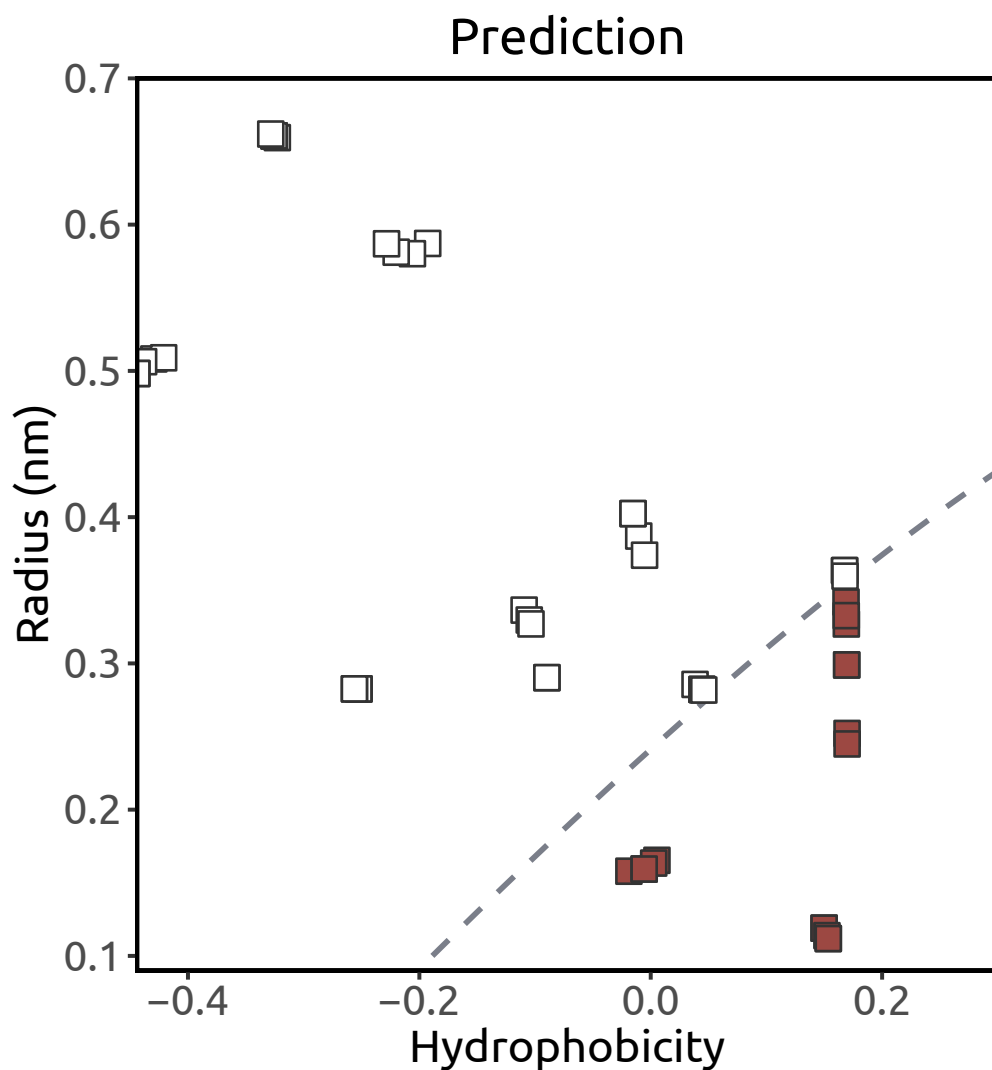
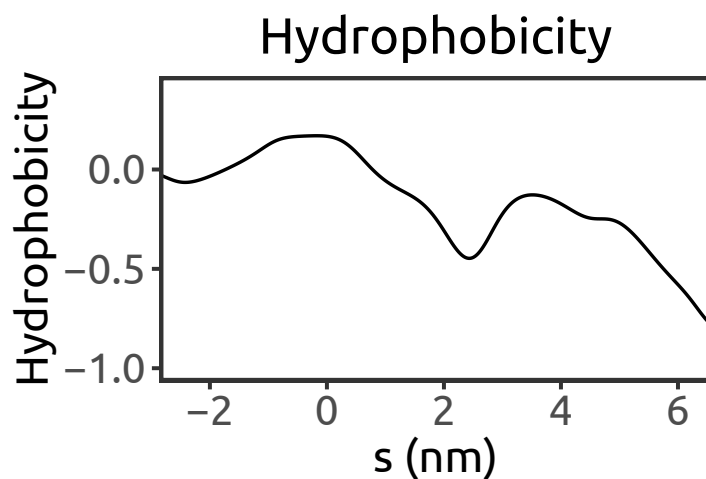
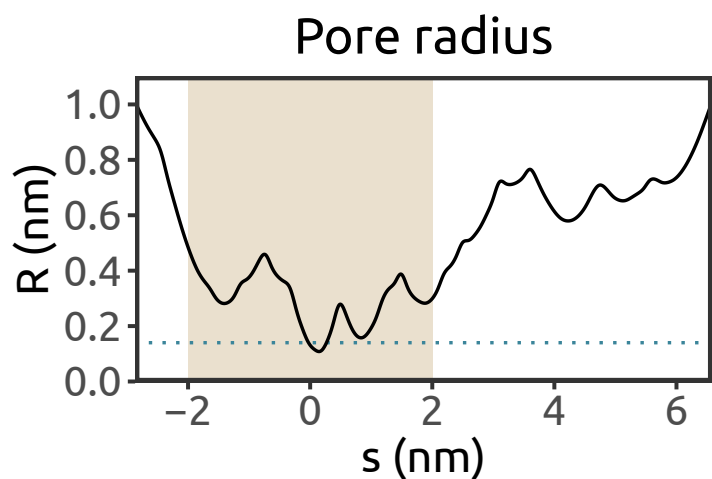
Simulation result:
barrier to water

GlyR (PDB ID: 3JAD)

Danio rerio

cryo-EM (3.9 Å)

Du et al., 2015



Heuristic score:
1.59 (n = 17)

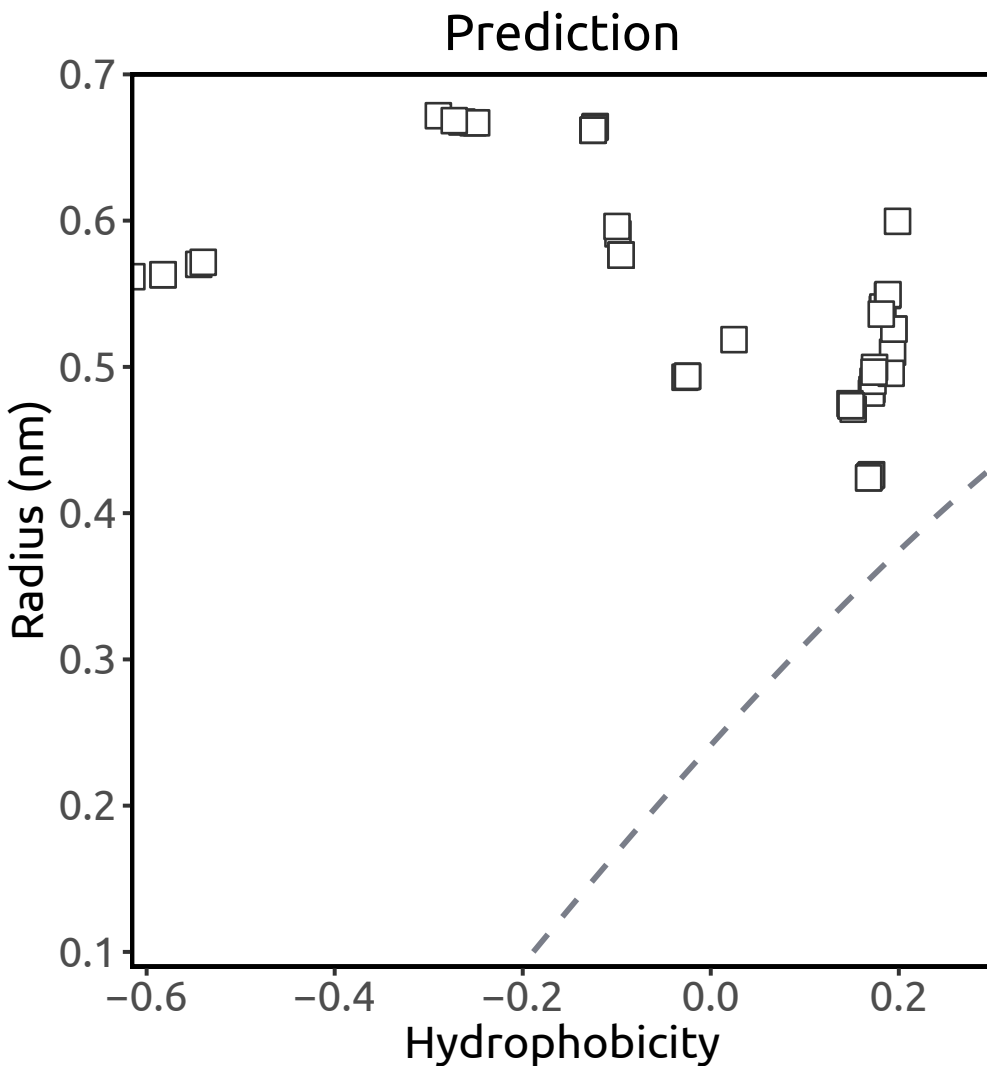
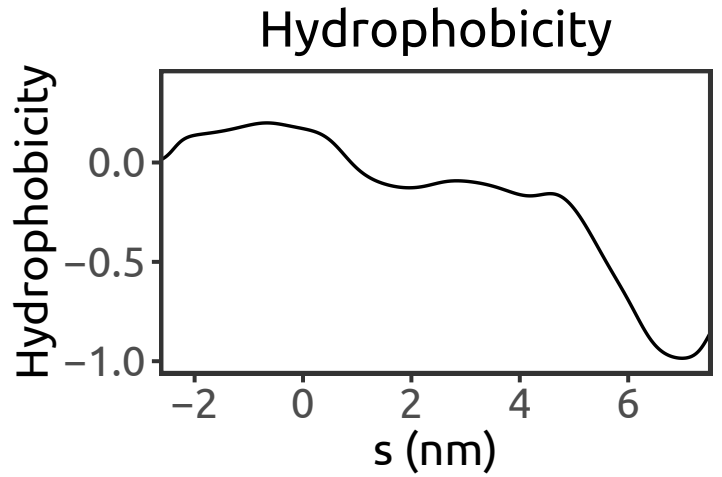
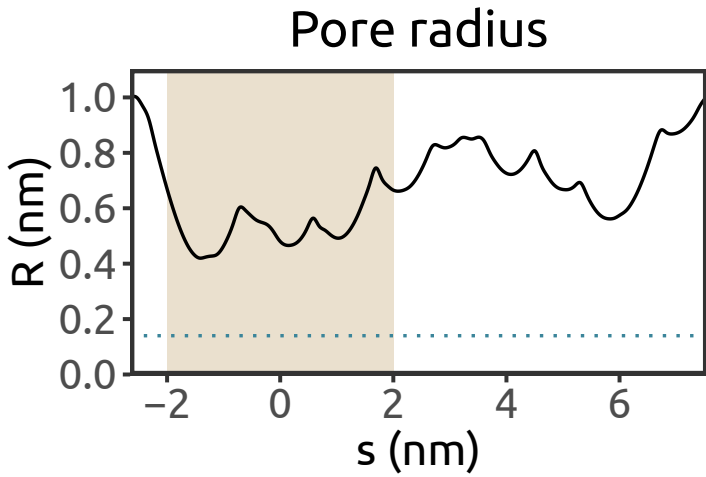
Simulation result:
barrier to water

GlyR (PDB ID: 3JAE)

Danio rerio

cryo-EM (3.9 Å)

Du et al., 2015



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

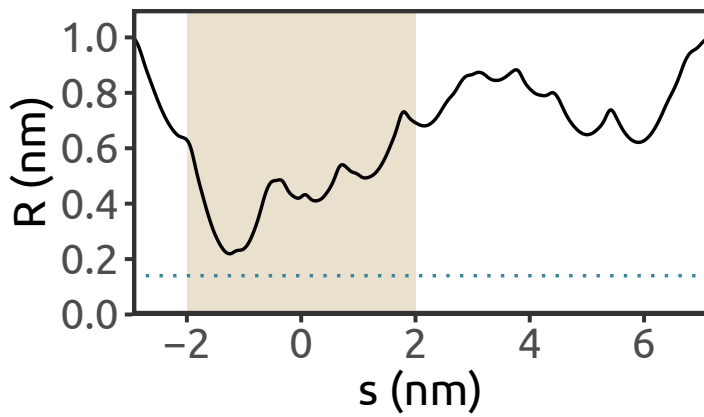
GlyR (PDB ID: 3JAF)

Danio rerio

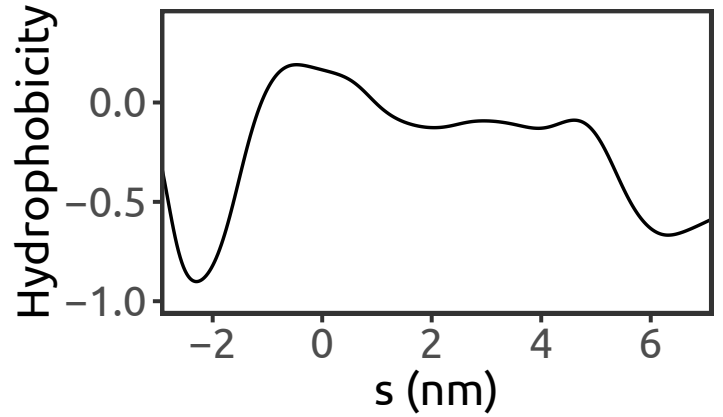
cryo-EM (3.8 Å)

Du et al., 2015

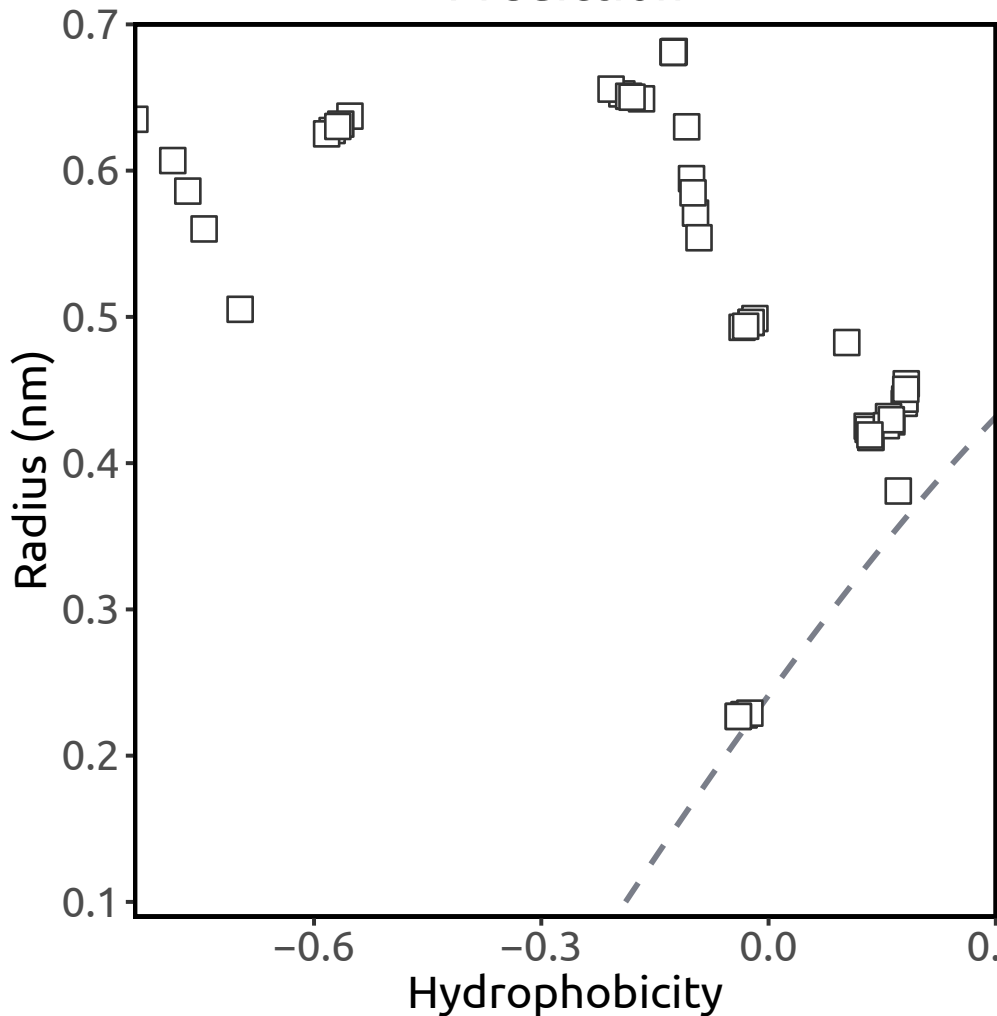
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 ($n = 0$)

Simulation result:
hydrated channel

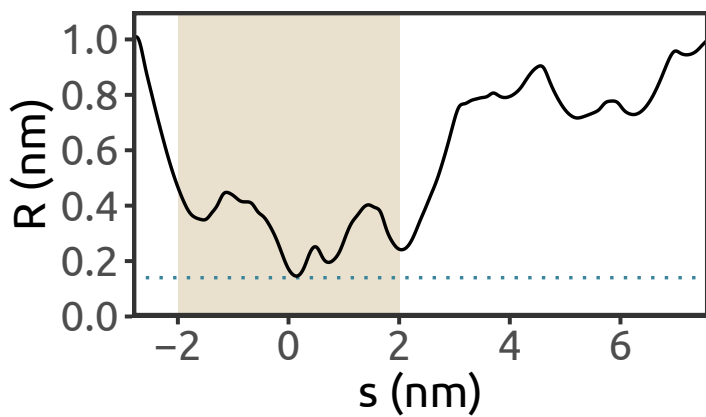
GlyR (PDB ID: 5CFB)

Homo sapiens

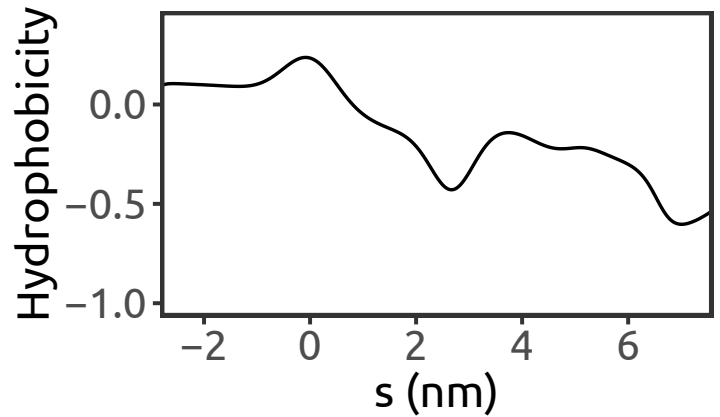
X-ray (3.04 Å)

Huang et al., 2015

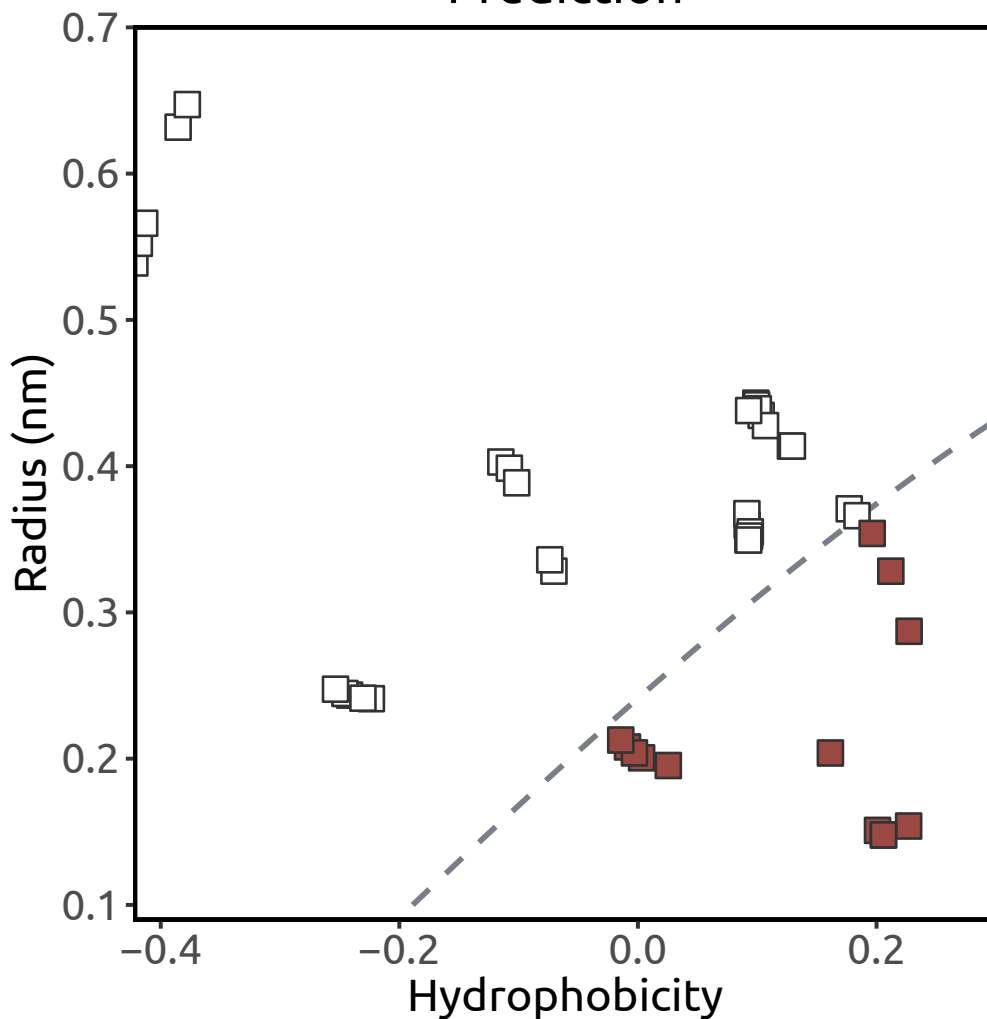
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.2 (n = 13)

Simulation result:
barrier to water

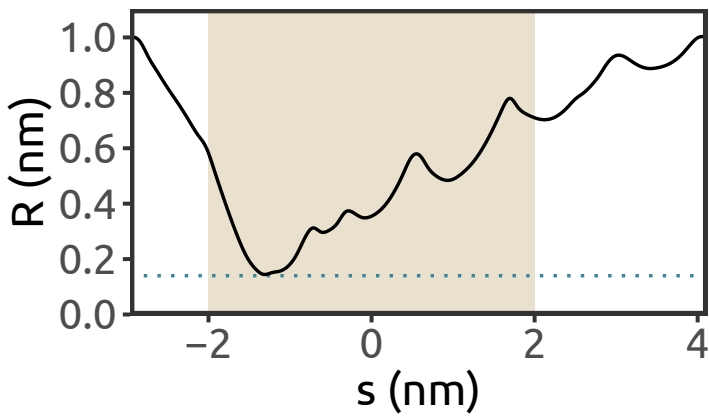
GlyR (PDB ID: 5VDH)

Homo sapiens

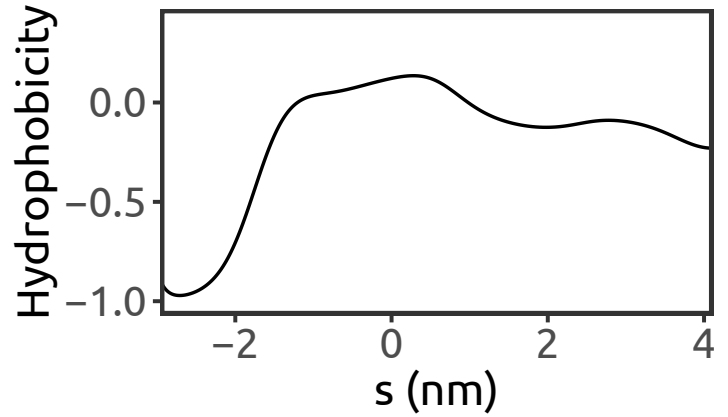
X-ray (2.85 Å)

Huang et al., 2017

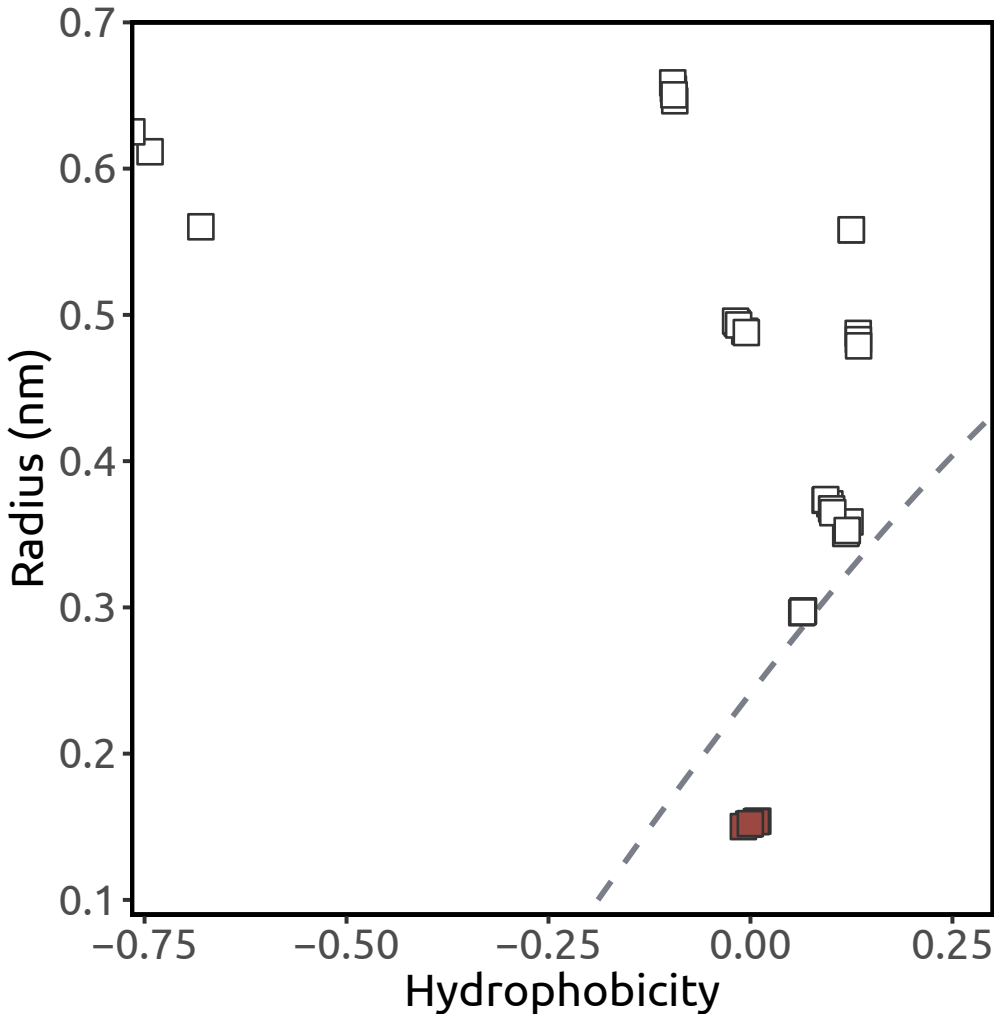
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.36 ($n = 5$)

Simulation result:
barrier to water

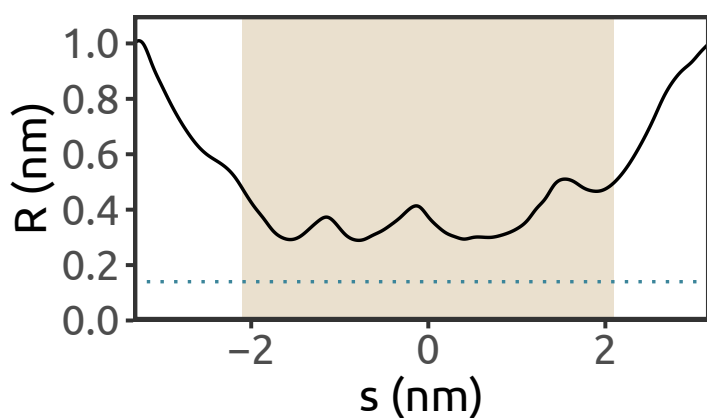
nAChR (PDB ID: 1OED)

Torpedo marmorata

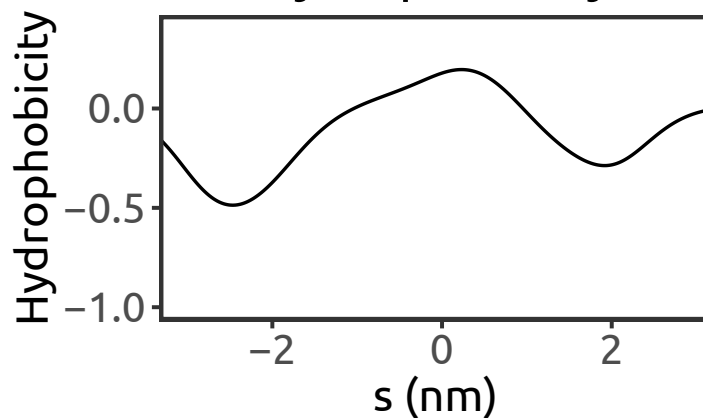
EM (4 Å)

Miyazawa et al., 2003

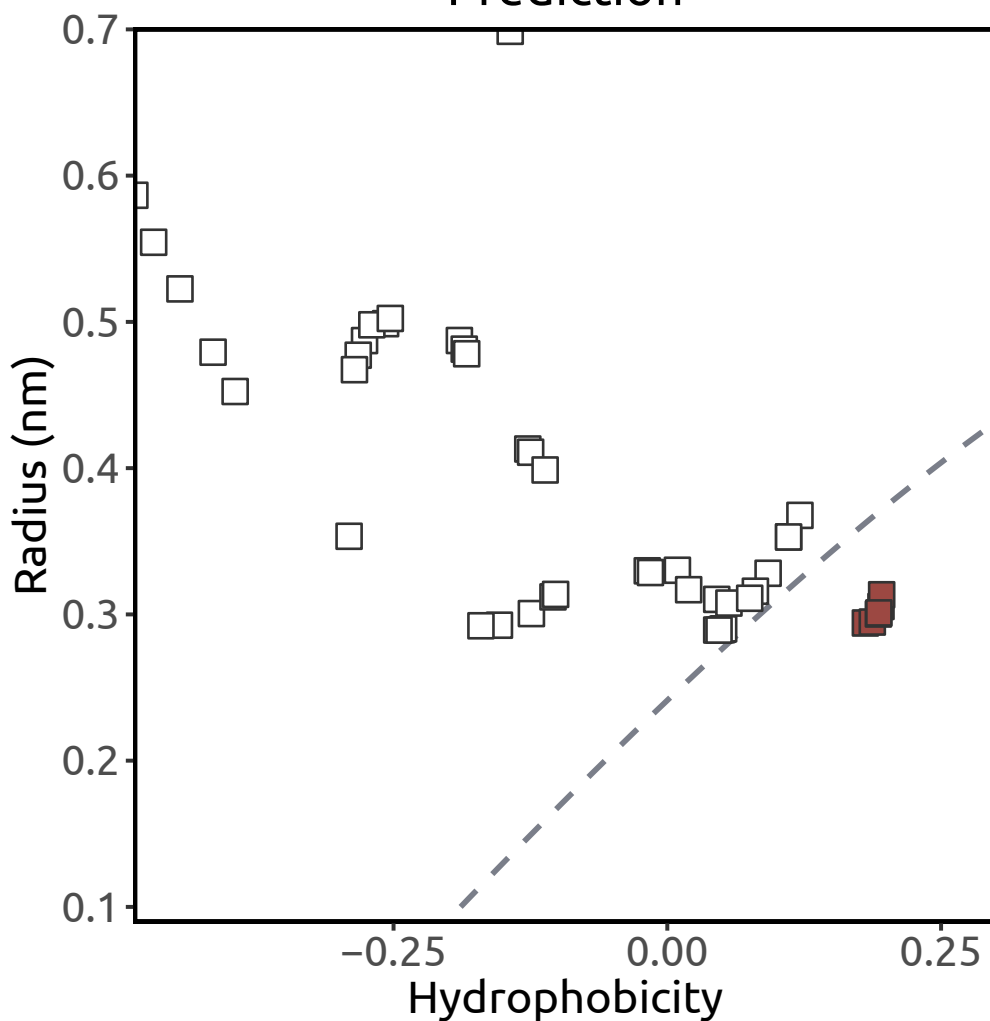
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.34 ($n = 6$)

Simulation result:
barrier to water

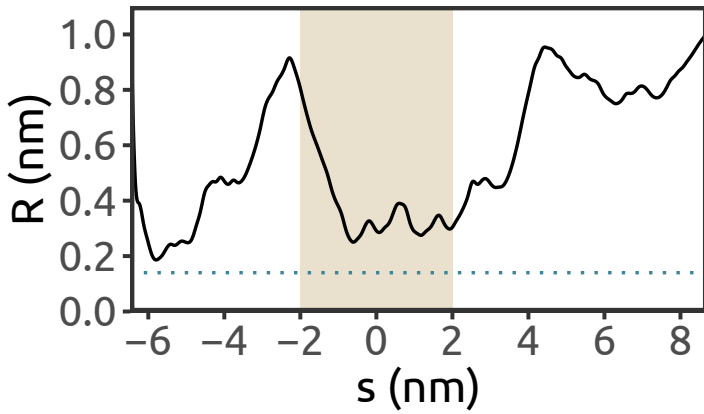
nAChR (PDB ID: 2BG9)

Torpedo marmorata

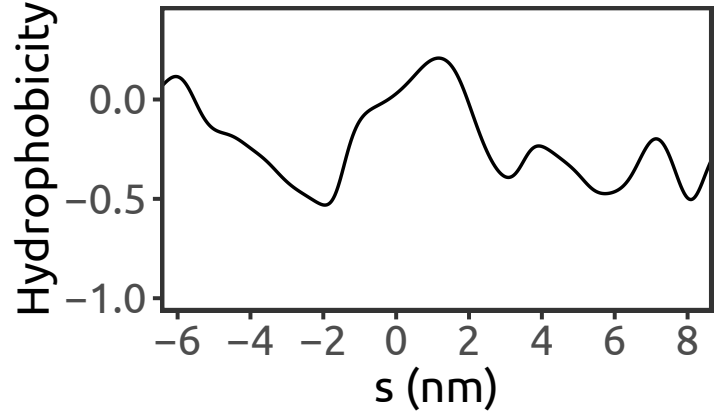
EM (4 Å)

Unwin, 2005

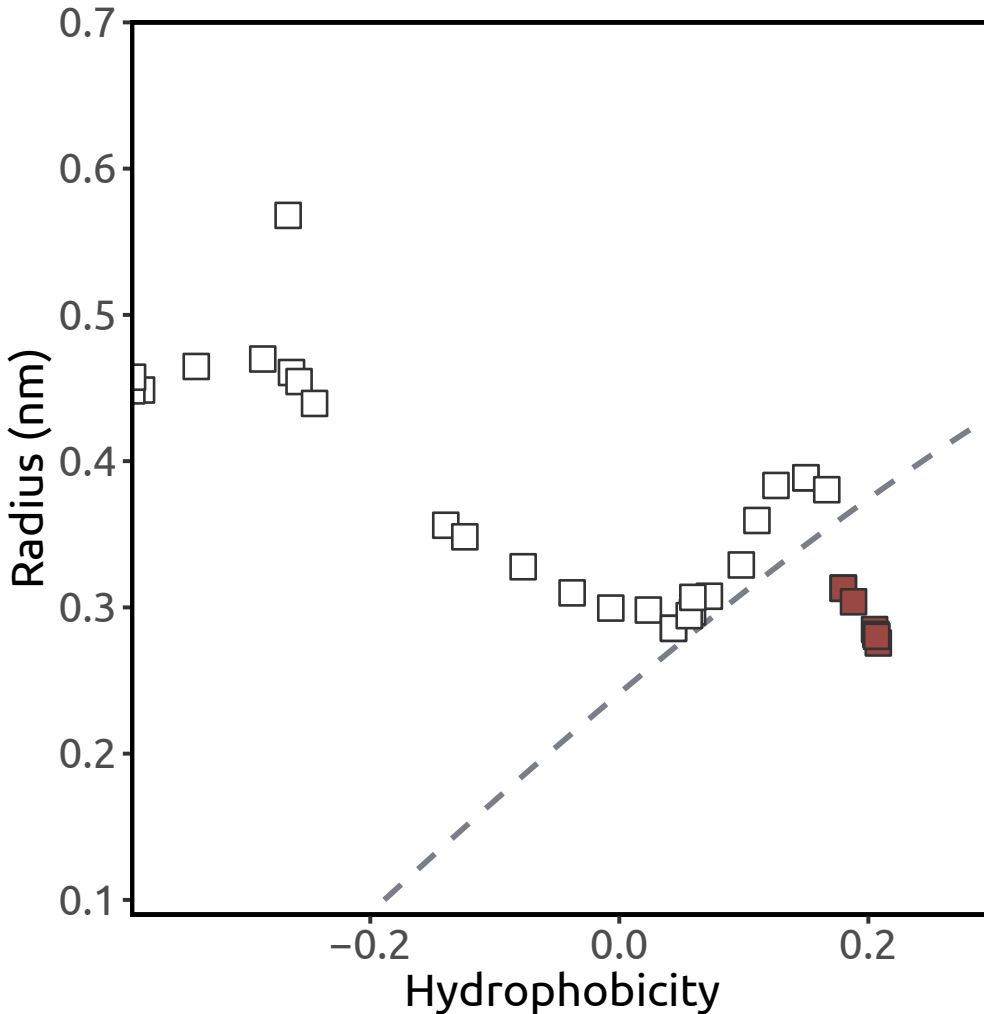
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.59 ($n = 9$)

Simulation result:

barrier to water

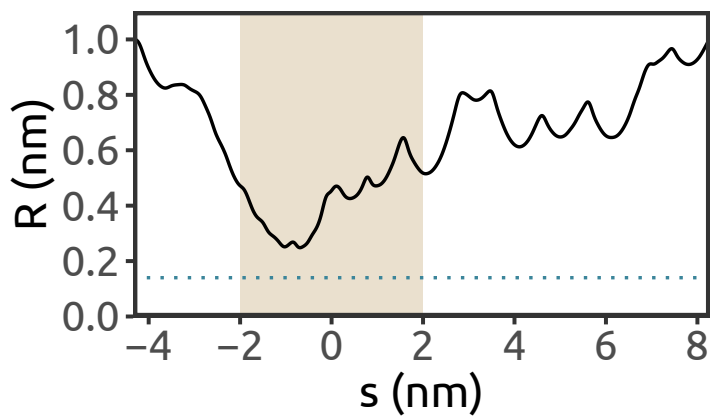
nAChR (PDB ID: 5KXI)

Homo sapiens

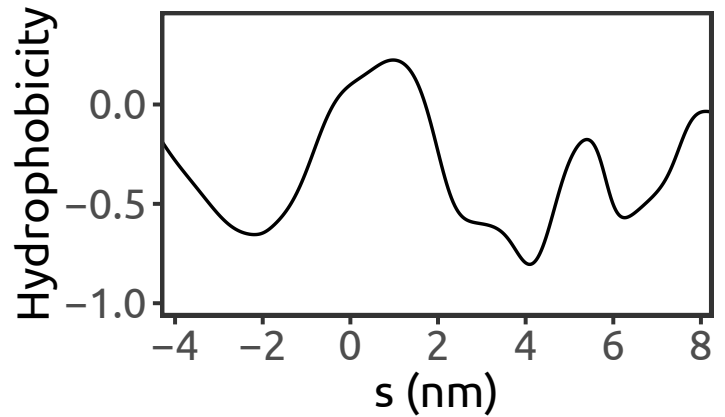
X-ray (3.94 Å)

Morales-Perez et al., 2016

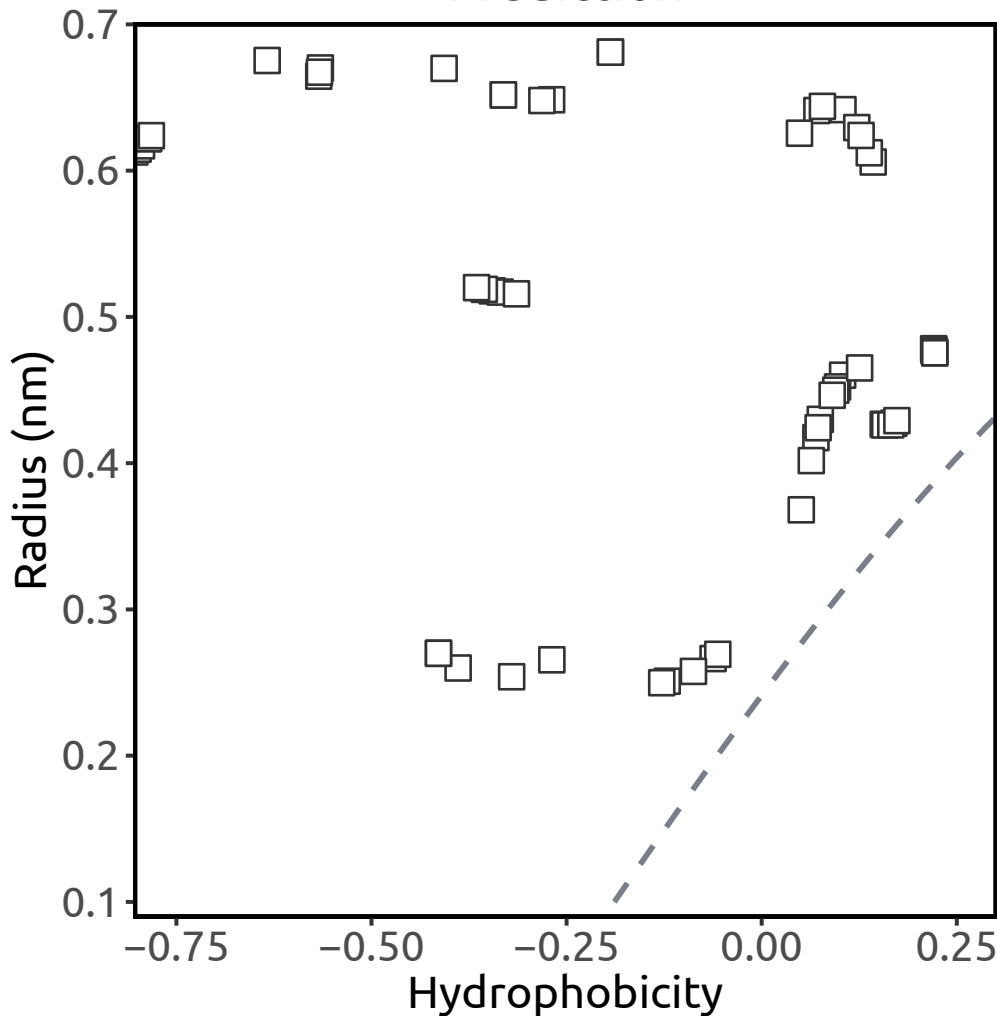
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

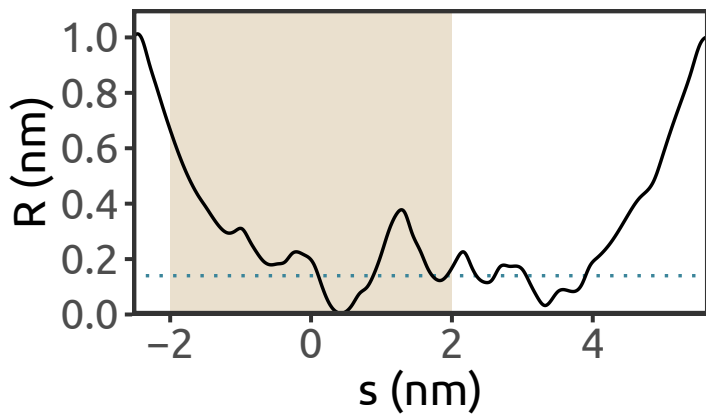
Simulation result:
hydrated channel

GluA (PDB ID: 5WEK)

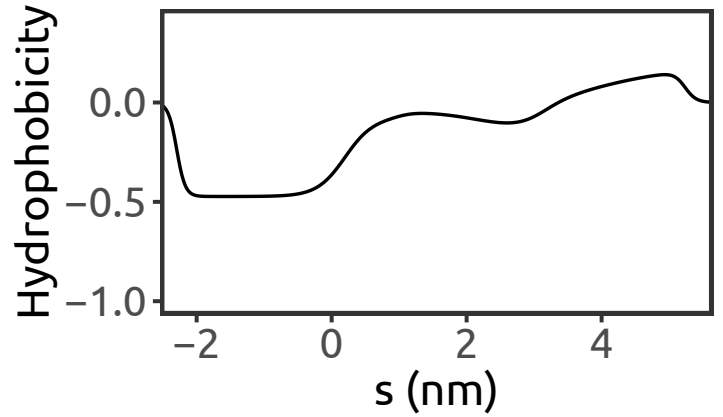
Rattus norvegicus
cryo-EM (4.6 Å)

Twomey et al., 2017

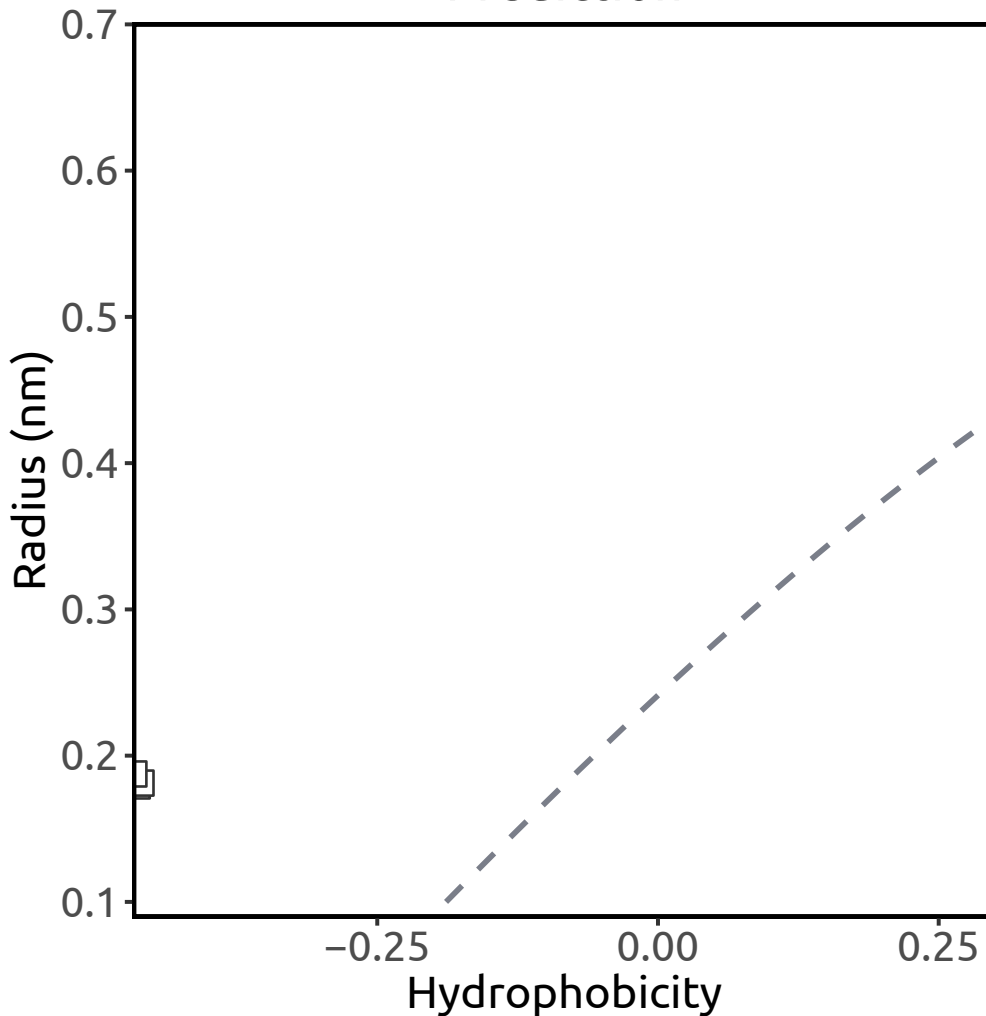
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.09 (n = 1)

Simulation result:
barrier to water

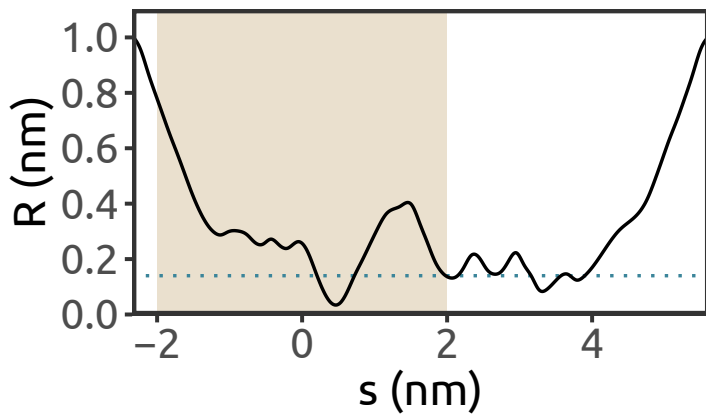
GluA (PDB ID: 5WEL)

Rattus norvegicus

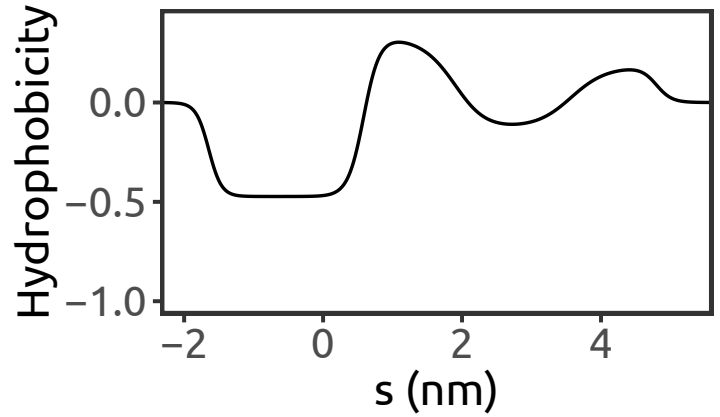
cryo-EM (4.4 Å)

Twomey et al., 2017

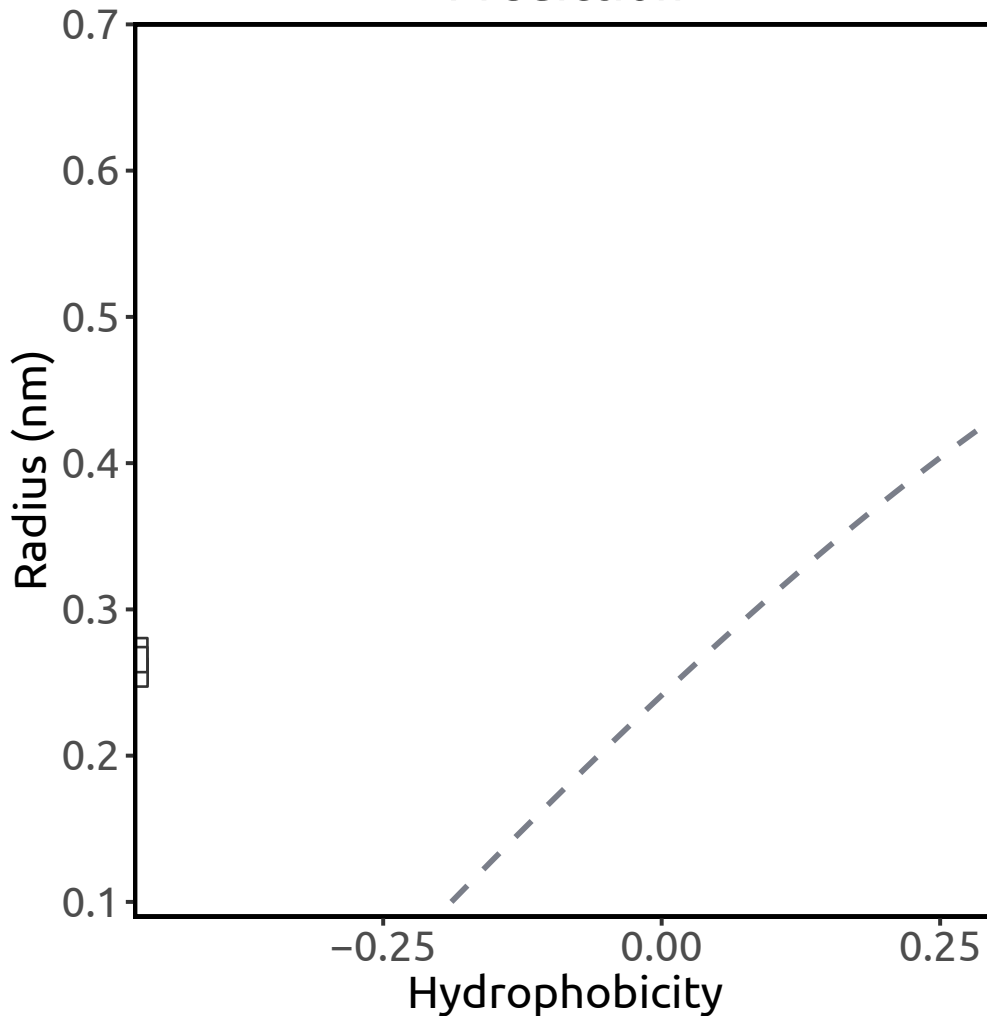
Pore radius



Hydrophobicity



Prediction



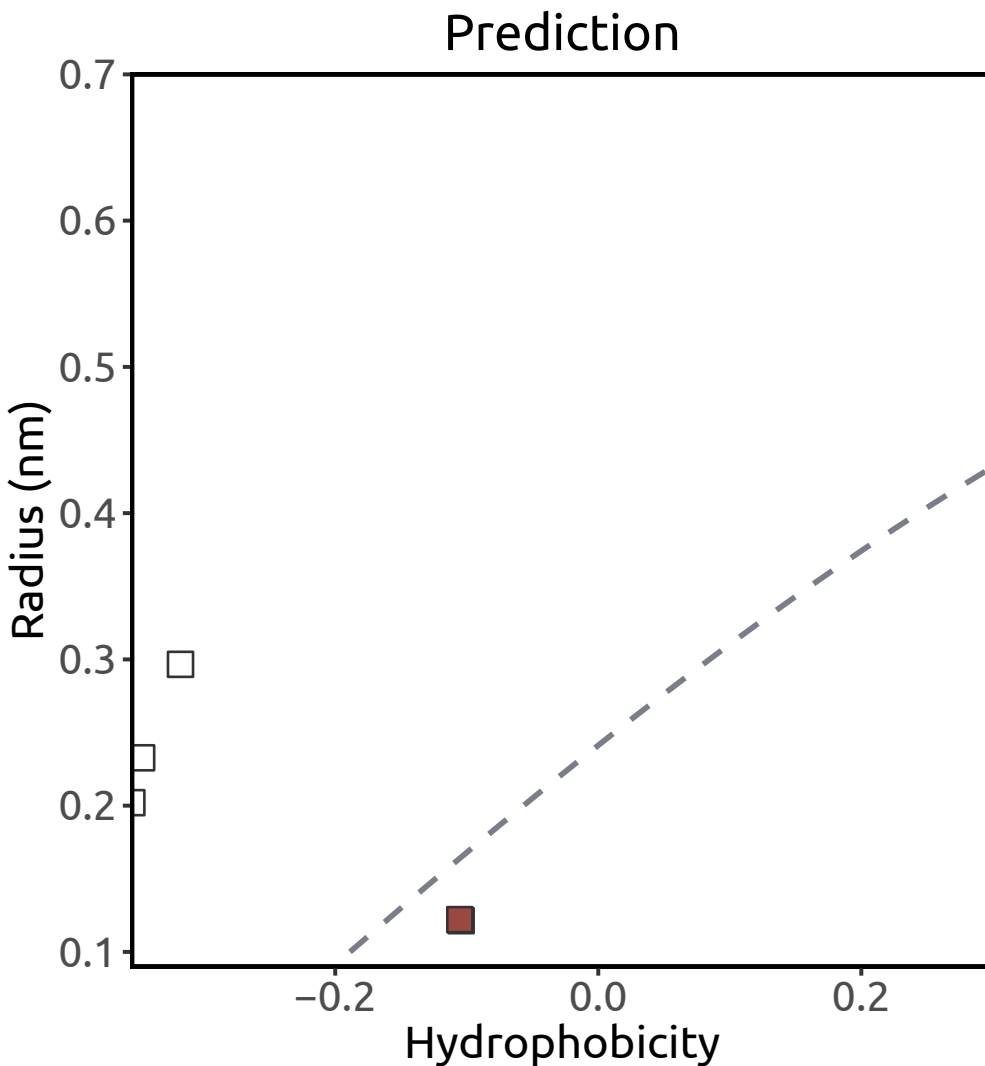
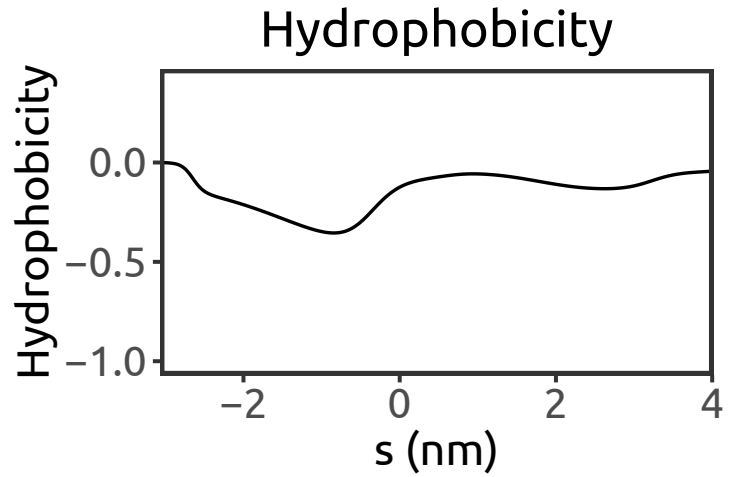
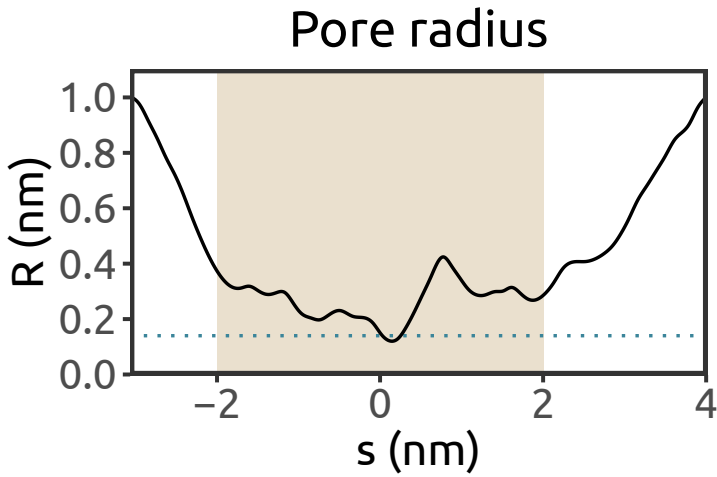
Heuristic score:
0 (n = 0)

Simulation result:
barrier to water

GluA (PDB ID: 5WEO)

Rattus norvegicus
cryo-EM (4.2 Å)

Twomey et al., 2017



Heuristic score:
0.07 (n = 2)

Simulation result:
hydrated channel

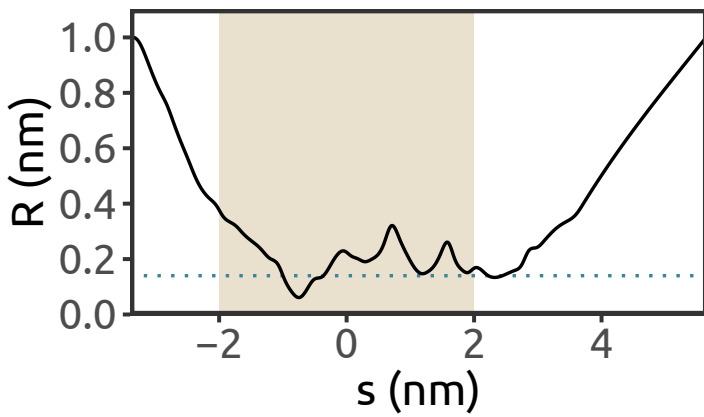
GluN (PDB ID: 5UOW)

Xenopus laevis

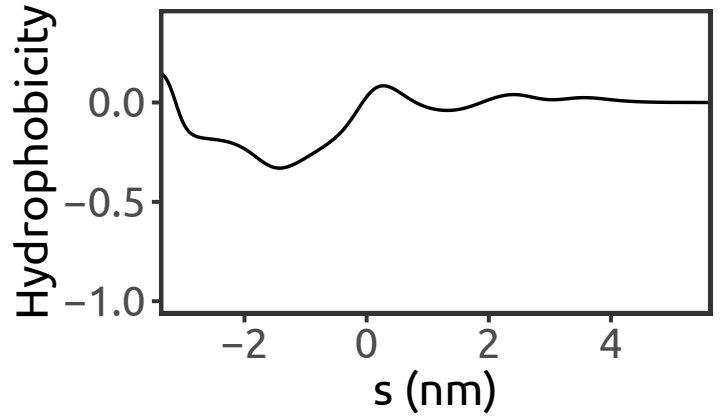
cryo-EM (4.5 Å)

Lü et al., 2017

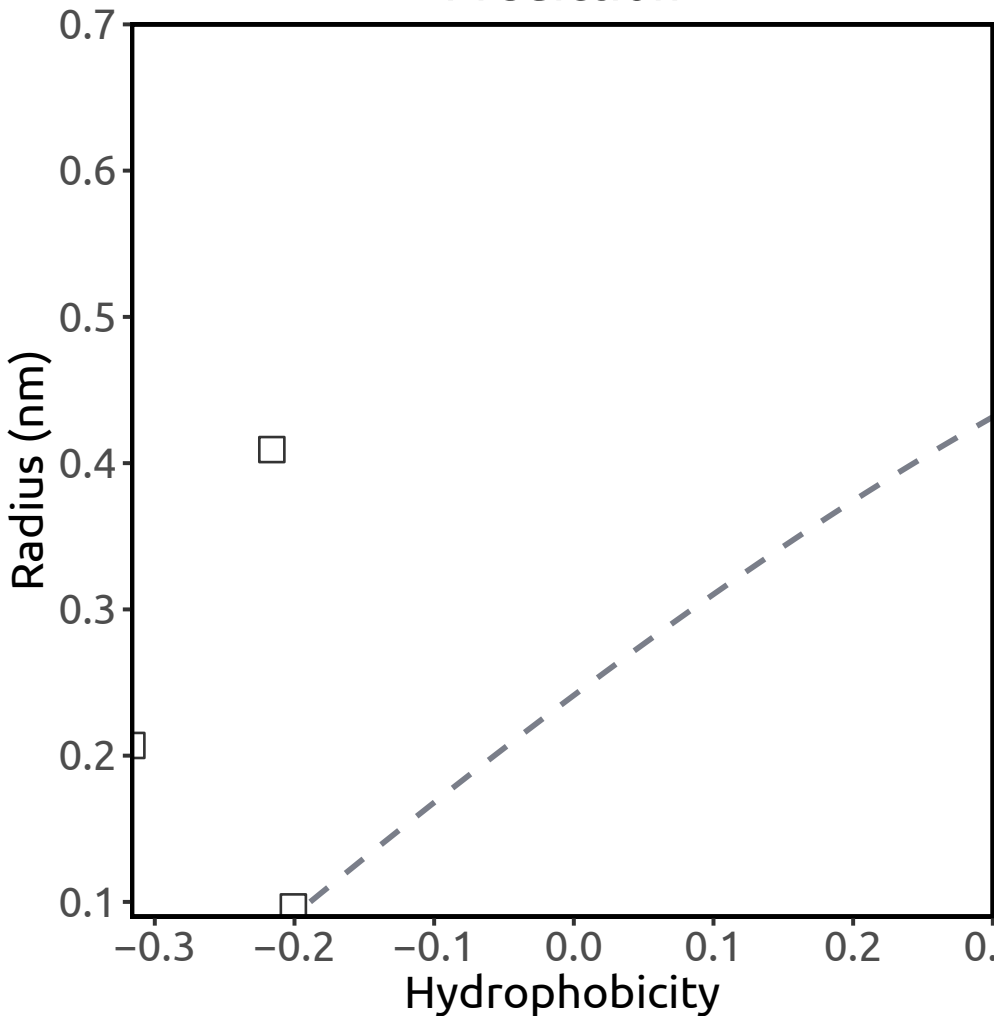
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

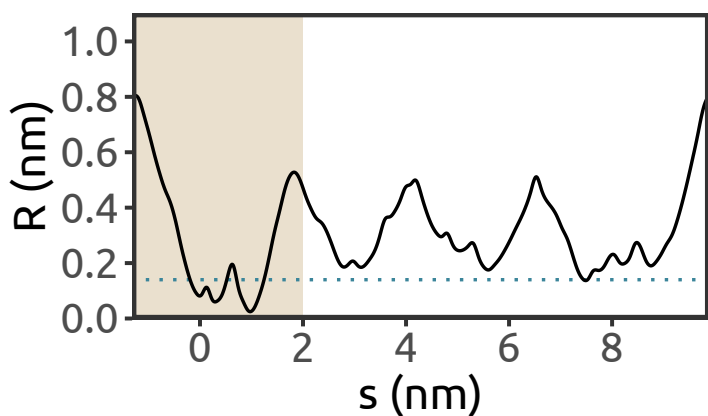
P2X3 (PDB ID: 5SVJ)

Homo sapiens

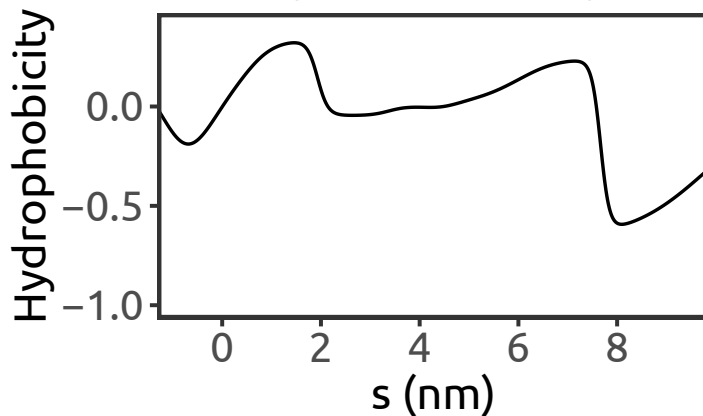
X-ray (2.98 Å)

Mansoor et al., 2016

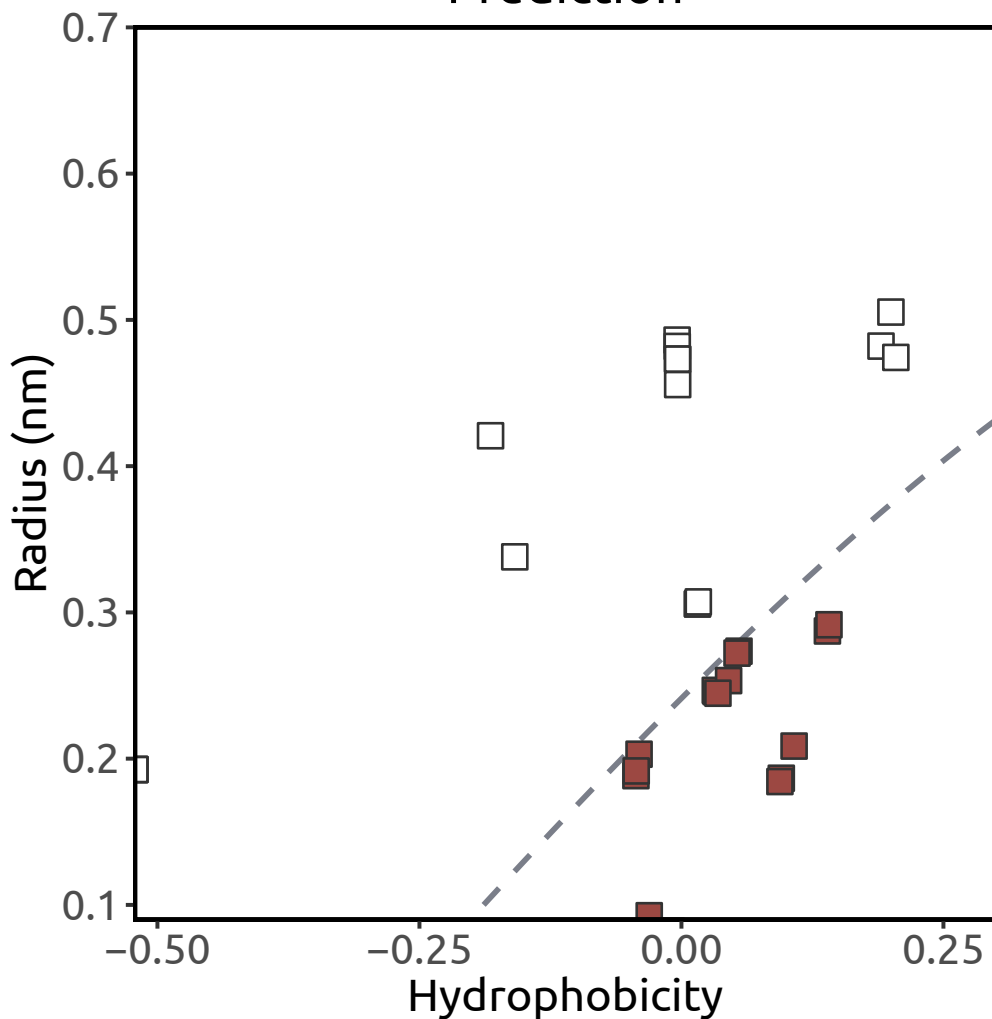
Pore radius



Hydrophobicity



Prediction



Heuristic score:

2.49 (n = 23)

Simulation result:

barrier to water

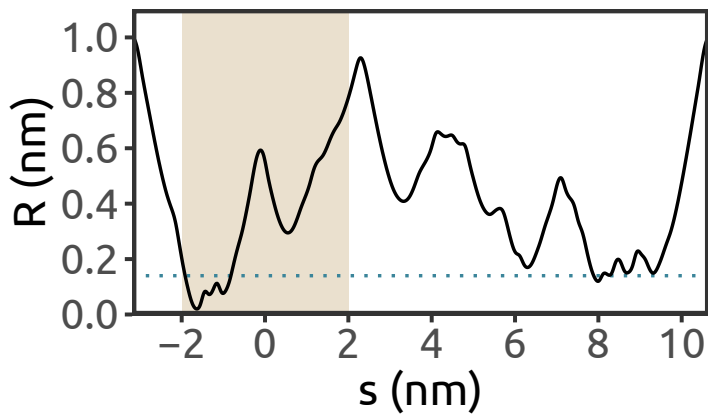
P2X3 (PDB ID: 5SVK)

Homo sapiens

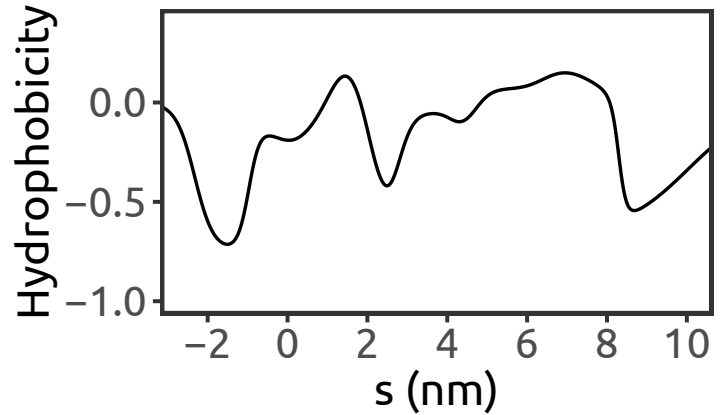
X-ray (2.77 Å)

Mansoor et al., 2016

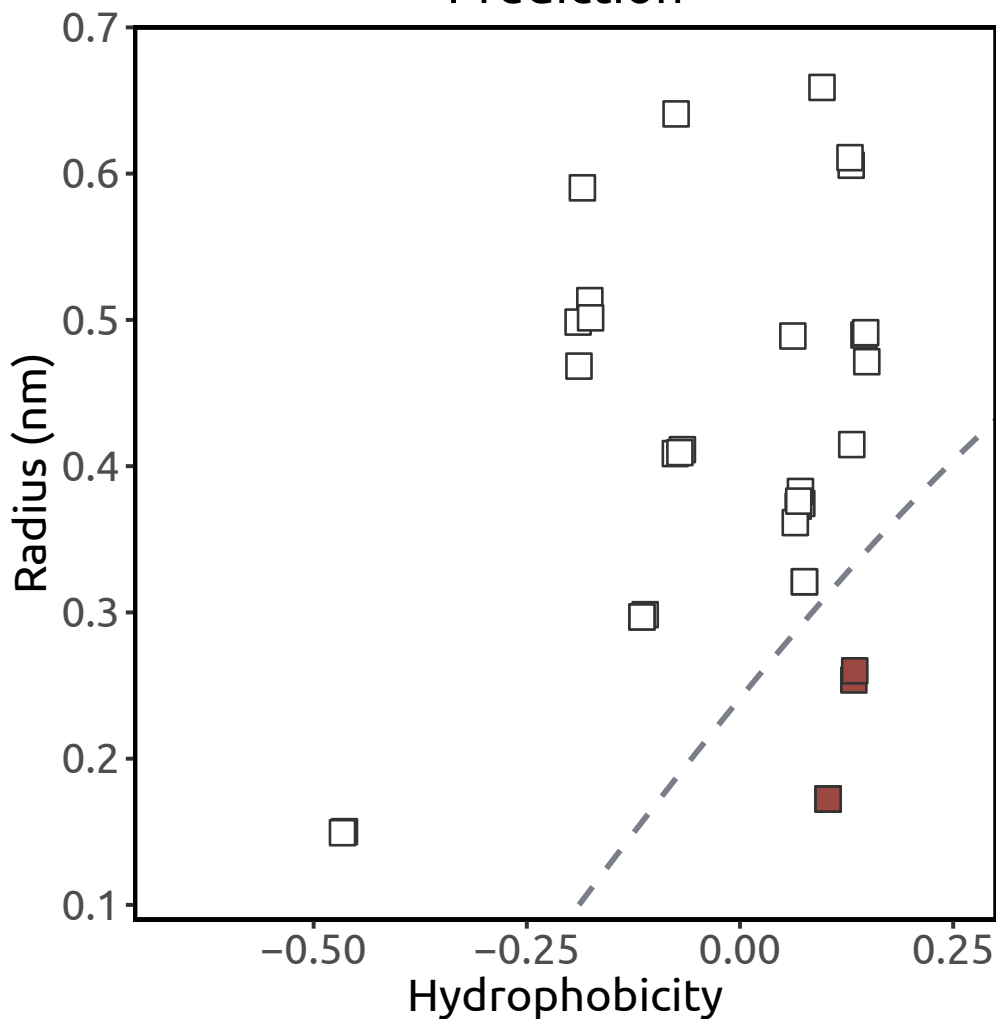
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.47 ($n = 5$)

Simulation result:
barrier to water

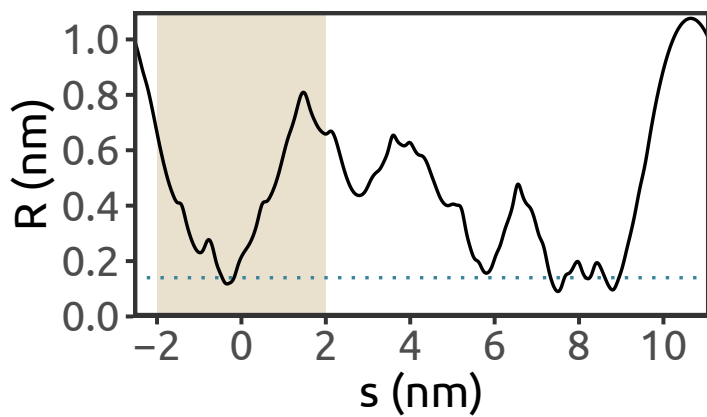
P2X3 (PDB ID: 5SVL)

Homo sapiens

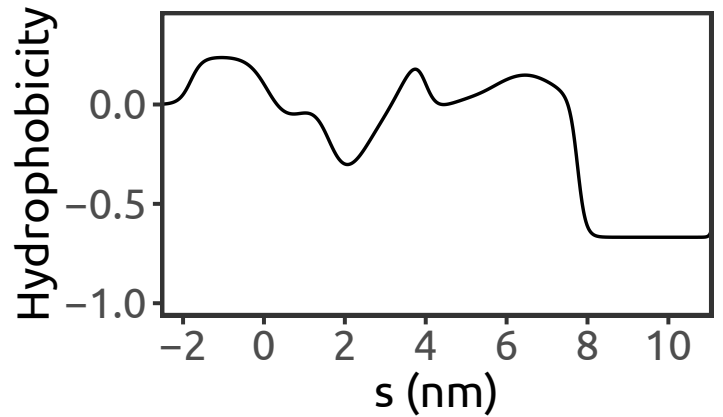
X-ray (2.9 Å)

Mansoor et al., 2016

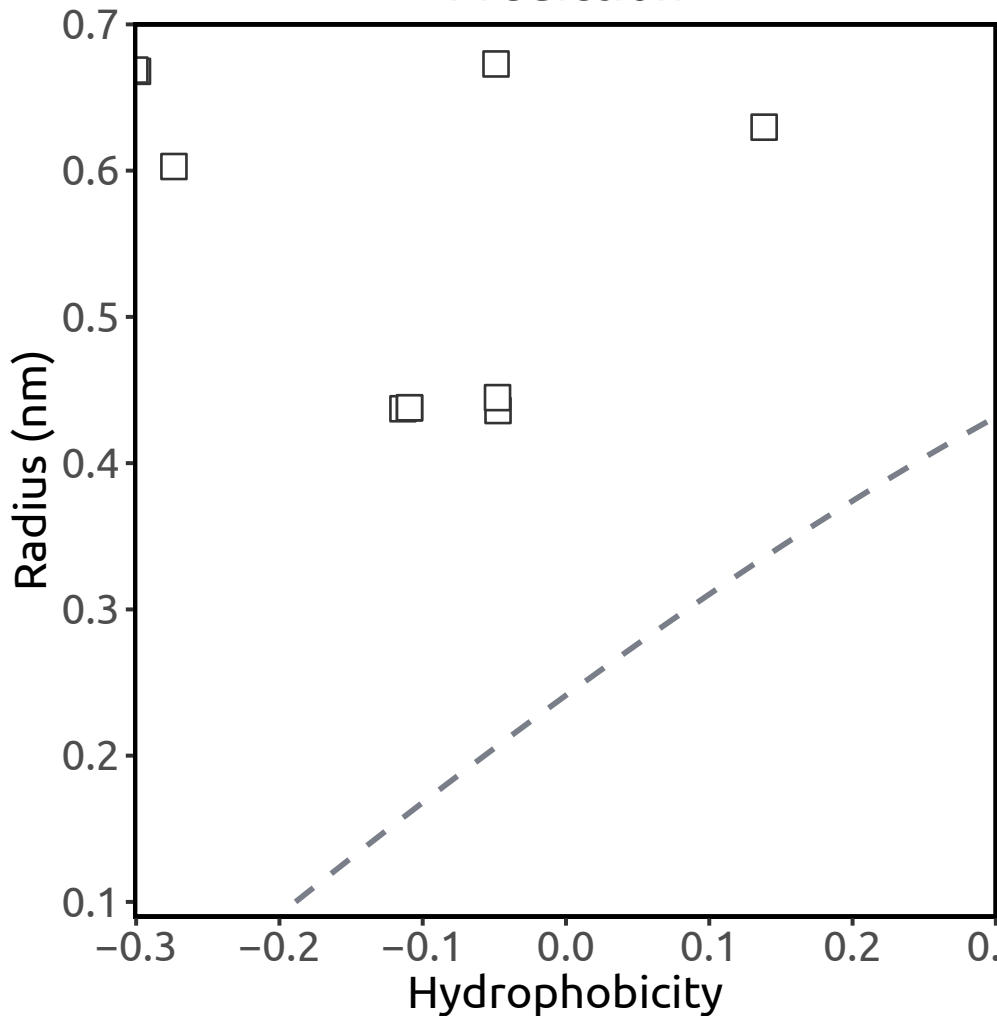
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

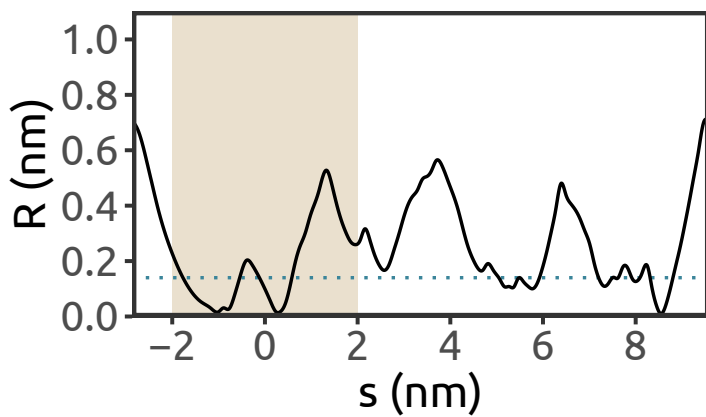
P2X4 (PDB ID: 3H9V)

Danio rerio

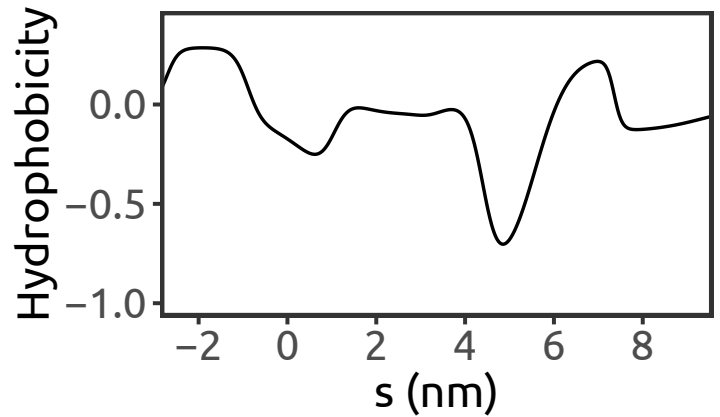
X-ray (3.1 Å)

Kawate et al., 2009

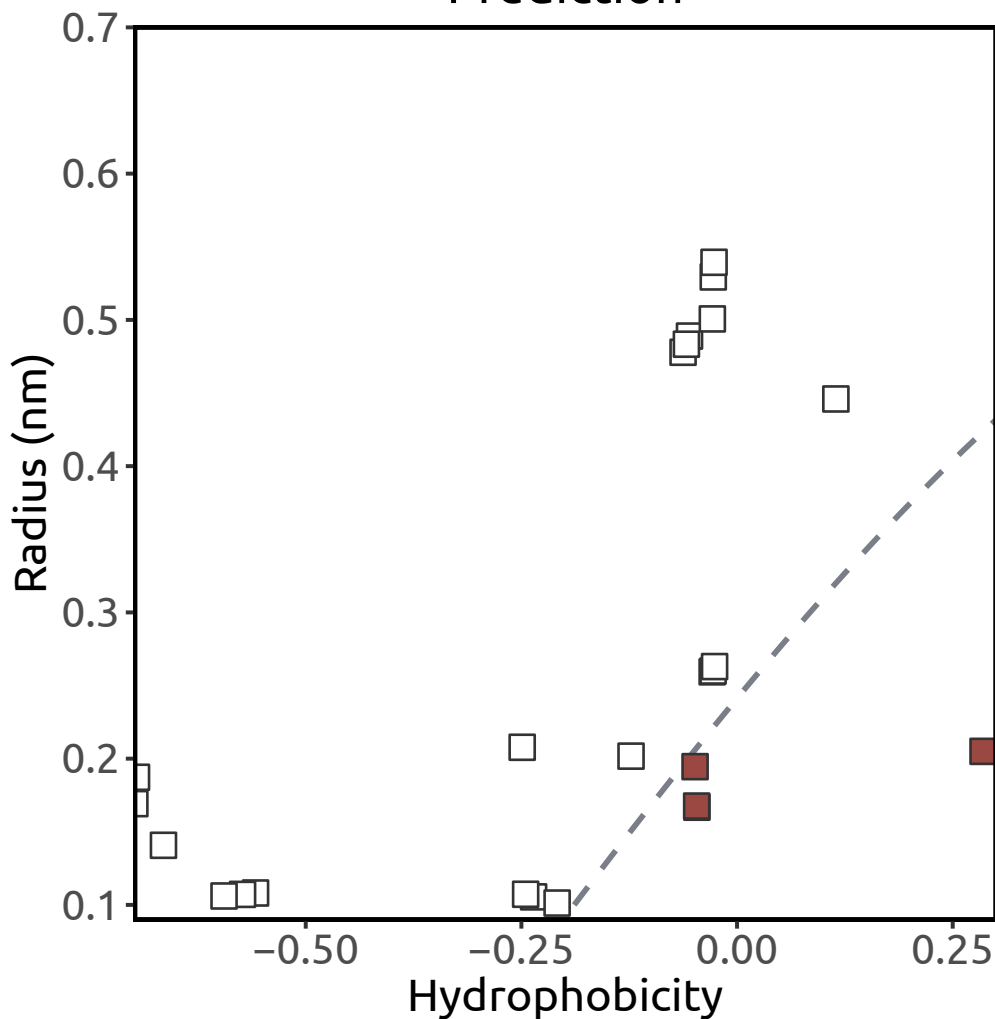
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.94 (n = 9)

Simulation result:
barrier to water

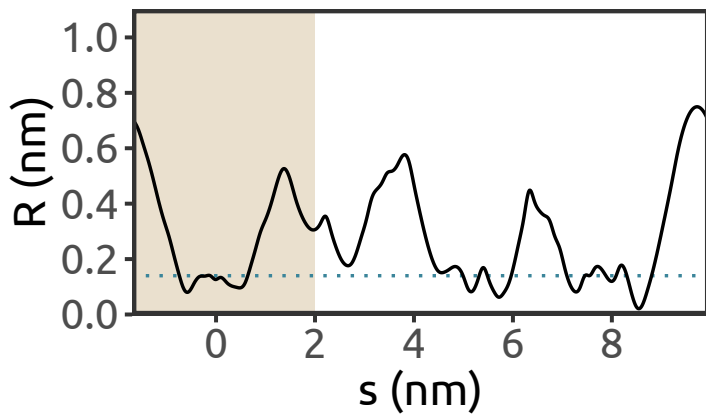
P2X4 (PDB ID: 3I5D)

Danio rerio

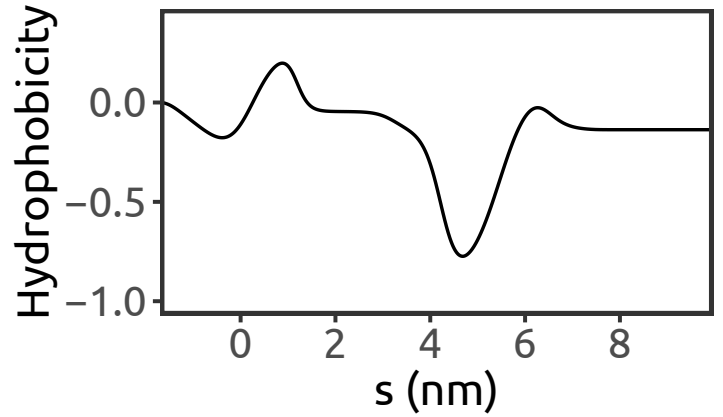
X-ray (3.46 Å)

Kawate et al., 2009

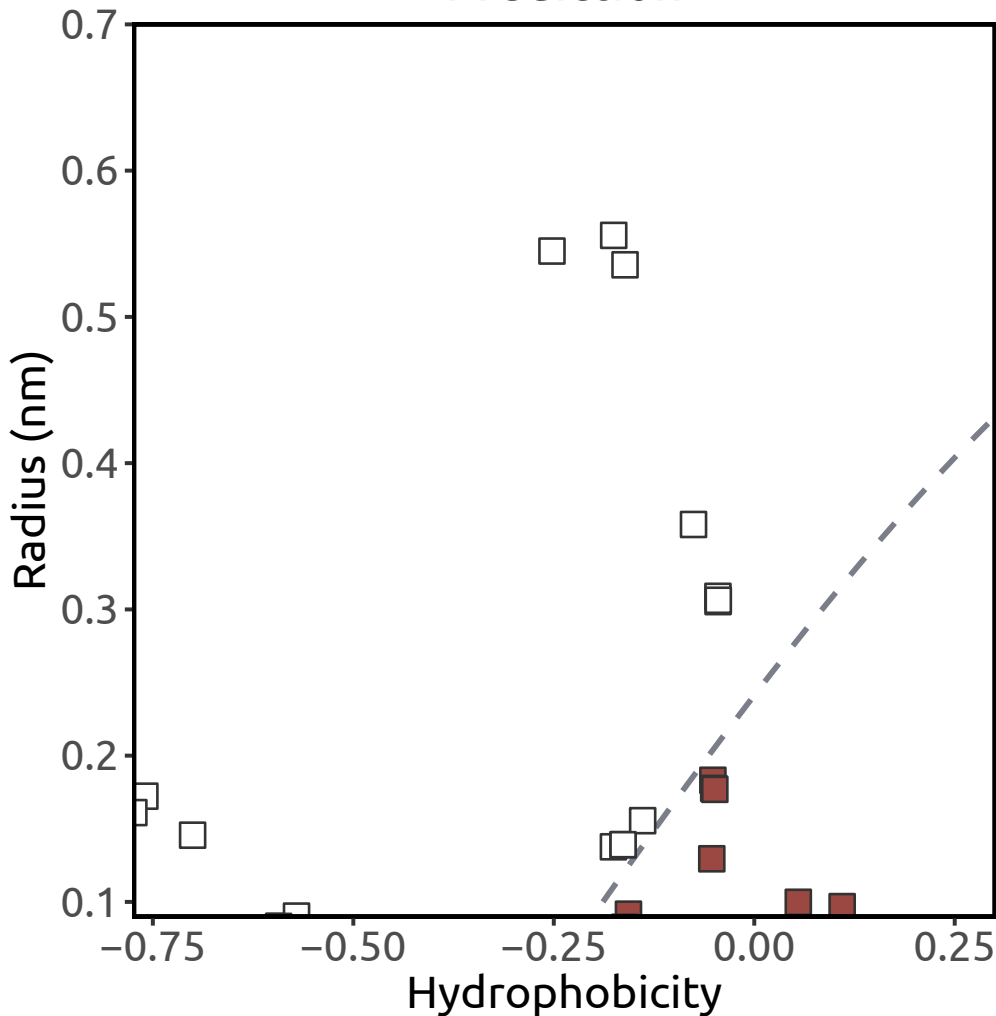
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.8 (n = 12)

Simulation result:
barrier to water

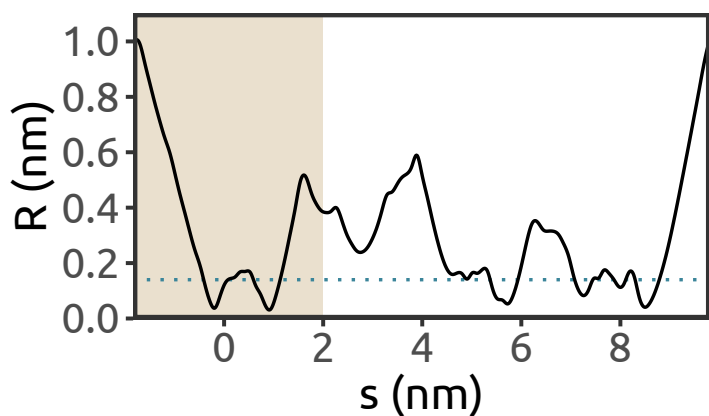
P2X4 (PDB ID: 4DW0)

Danio rerio

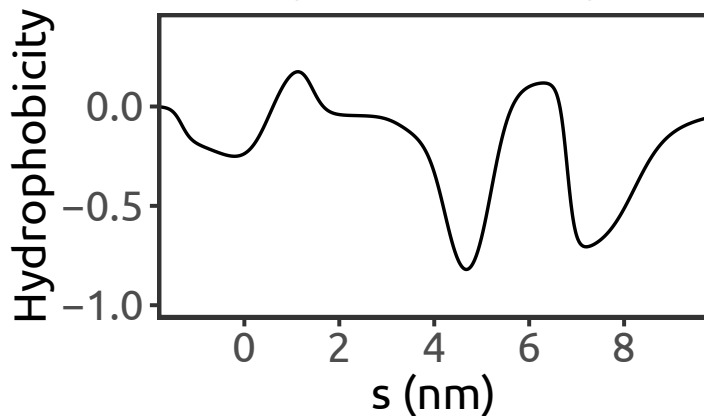
X-ray (2.9 Å)

Hattori & Gouaux, 2012

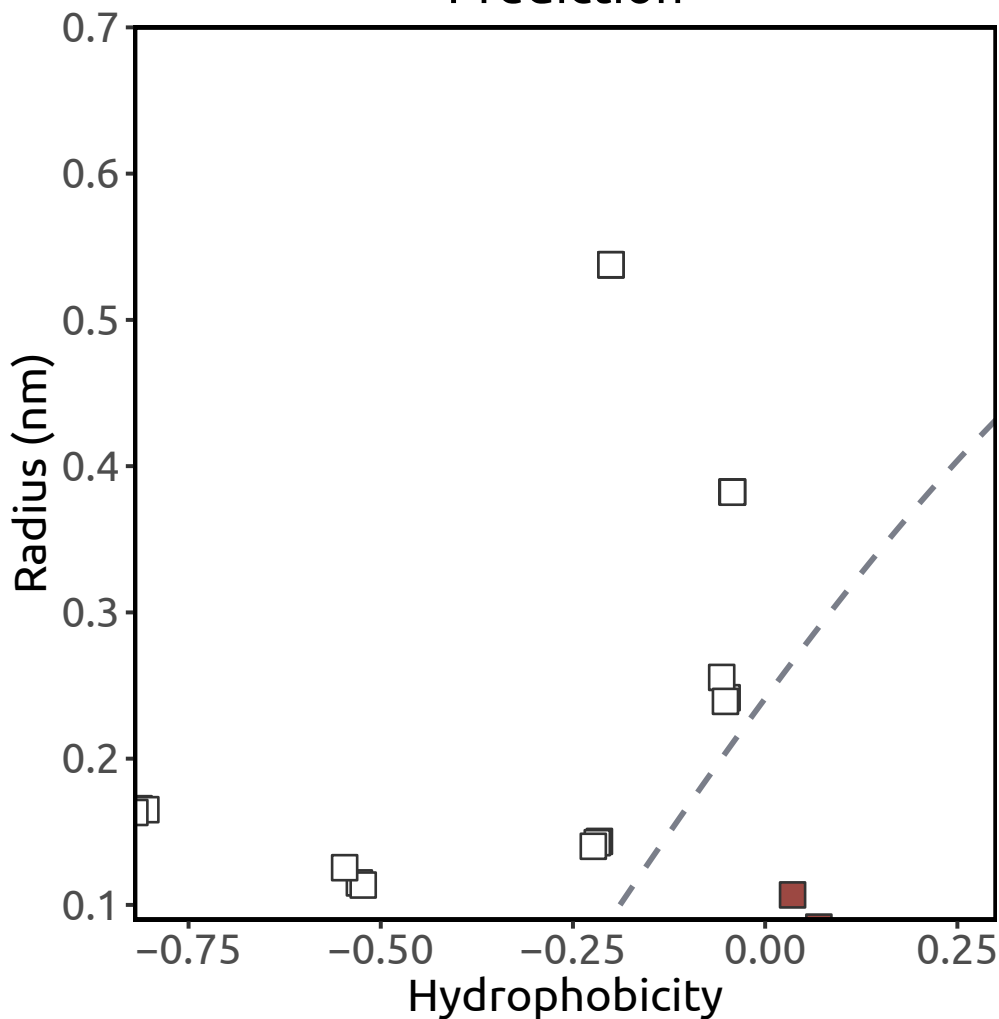
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.92 (n = 6)

Simulation result:

barrier to water

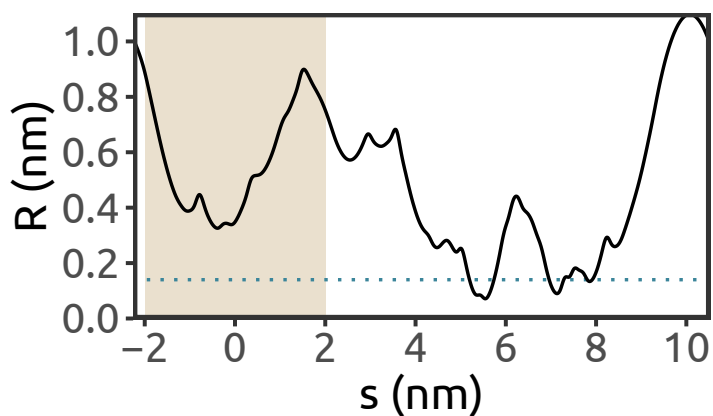
P2X4 (PDB ID: 4DW1)

Danio rerio

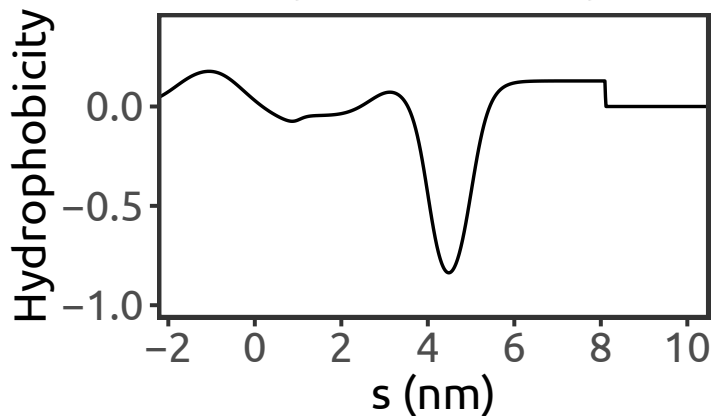
X-ray (2.8 Å)

Hattori & Gouaux, 2012

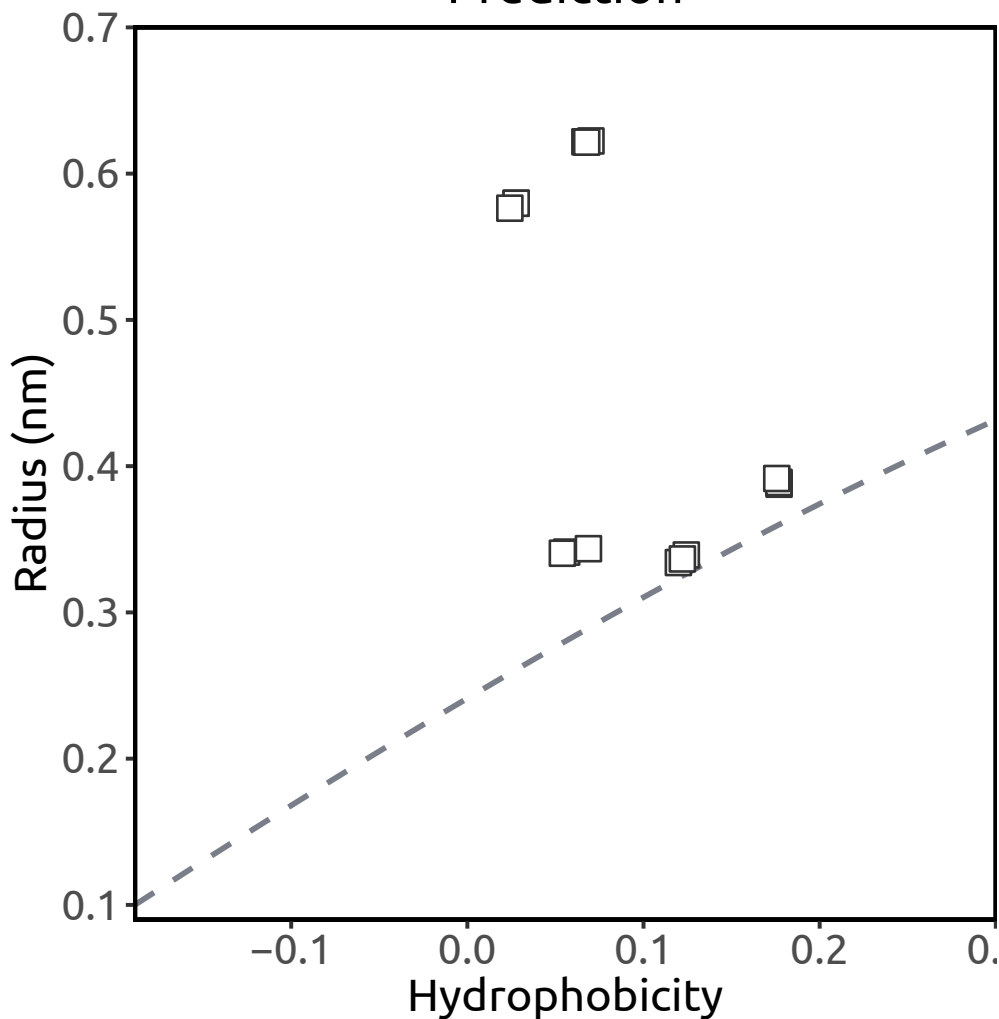
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

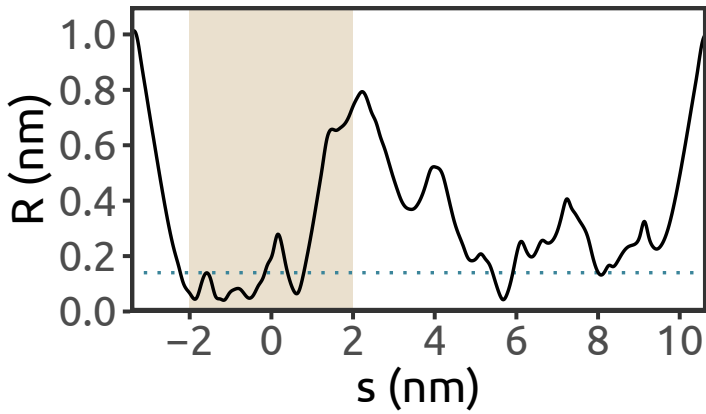
P2X7 (PDB ID: 5XW6)

Gallus gallus

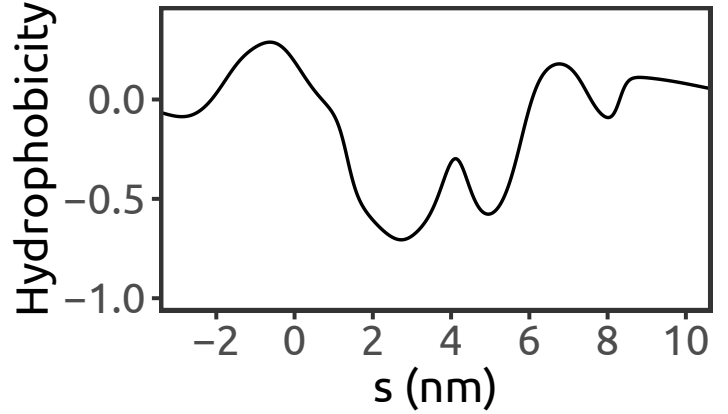
X-ray (3.1 Å)

Kasuya et al., 2017

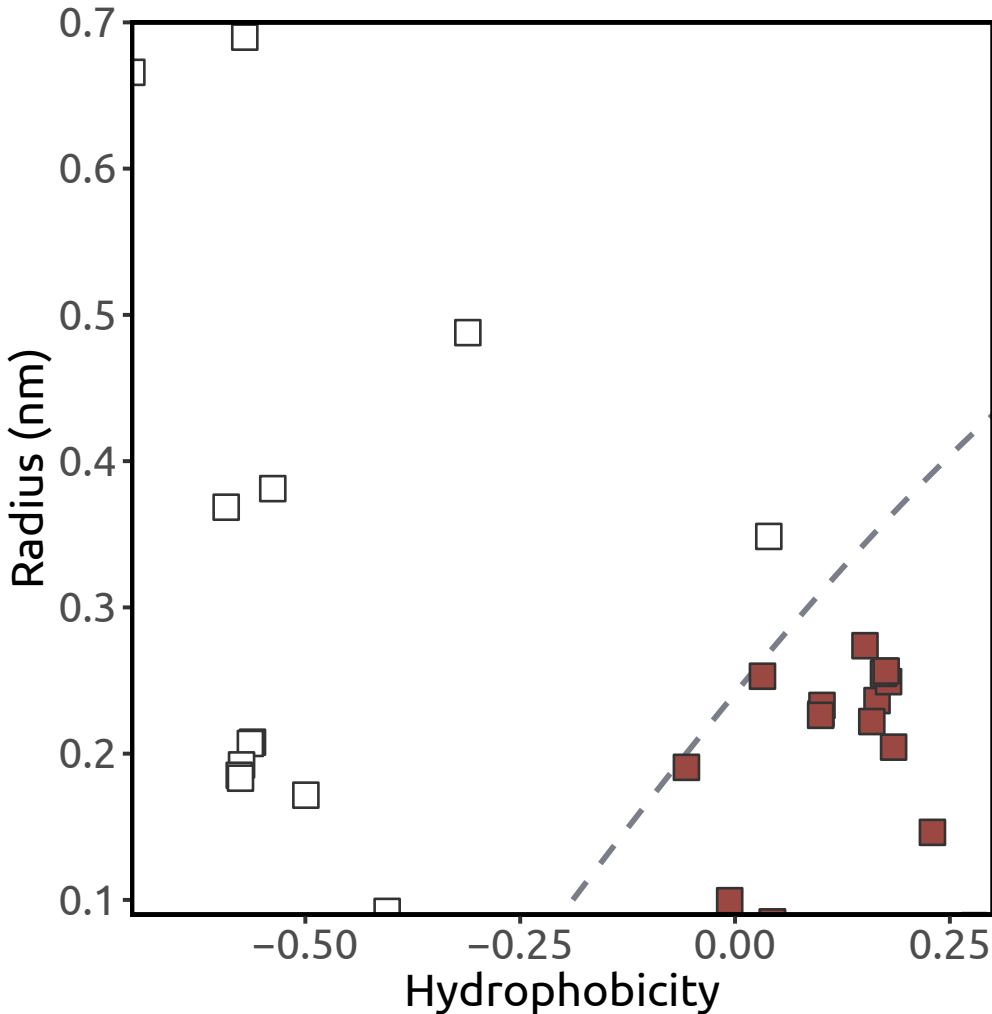
Pore radius



Hydrophobicity



Prediction



Heuristic score:
3.62 (n = 25)

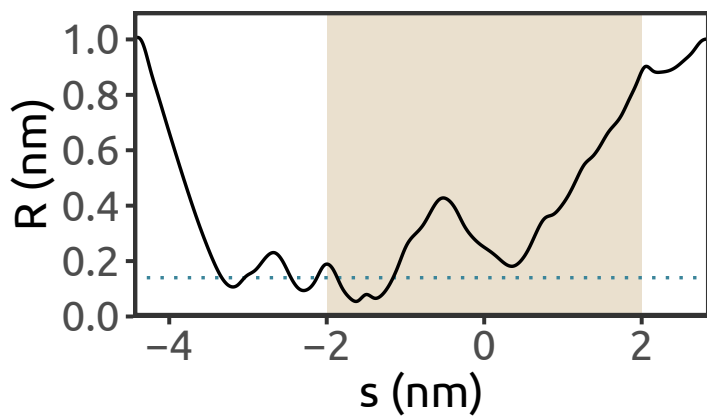
Simulation result:
barrier to water

NOMPC (PDB ID: 5VKQ)

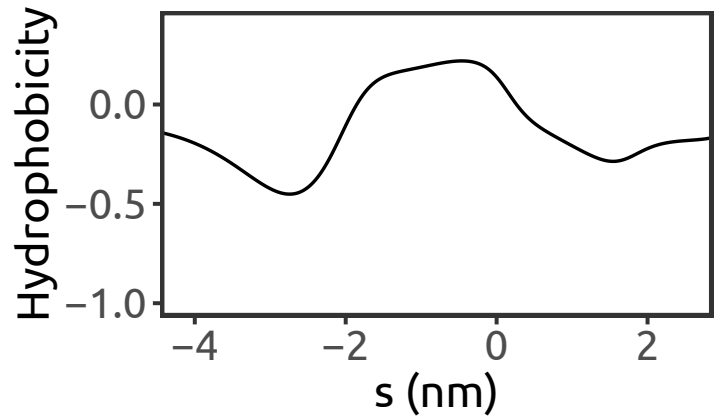
Drosophila melanogaster
cryo-EM (3.55 Å)

Jin et al., 2017

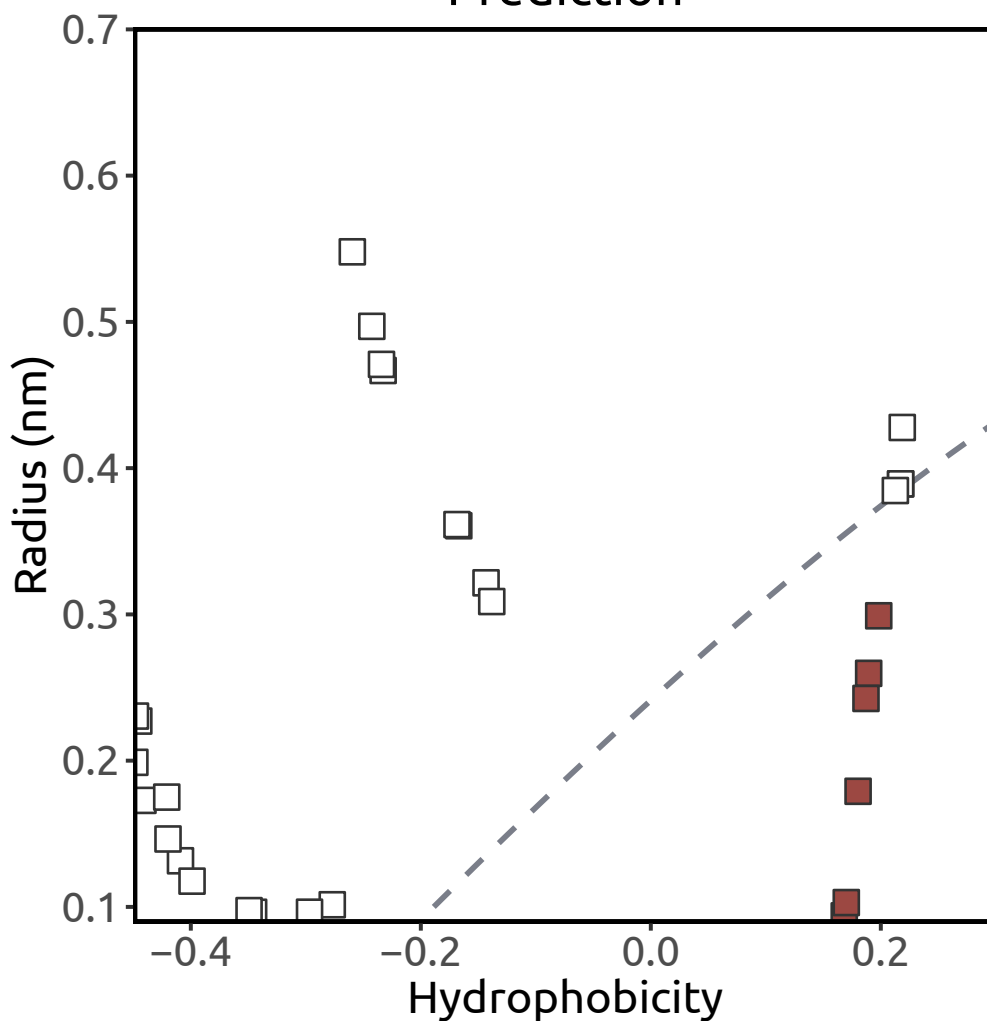
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.14 (n = 12)

Simulation result:
barrier to water

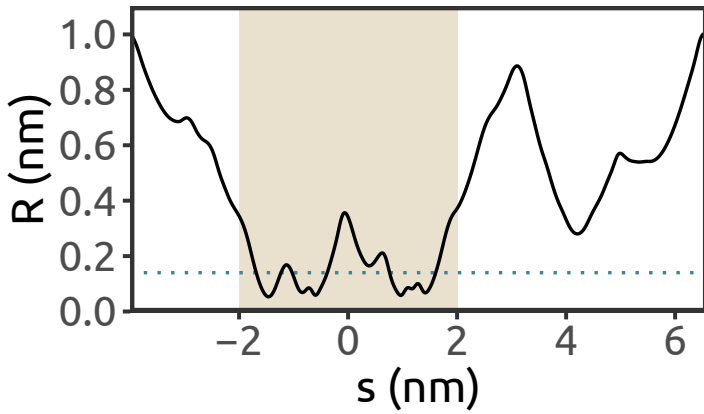
PKD2 (PDB ID: 5K47)

Homo sapiens

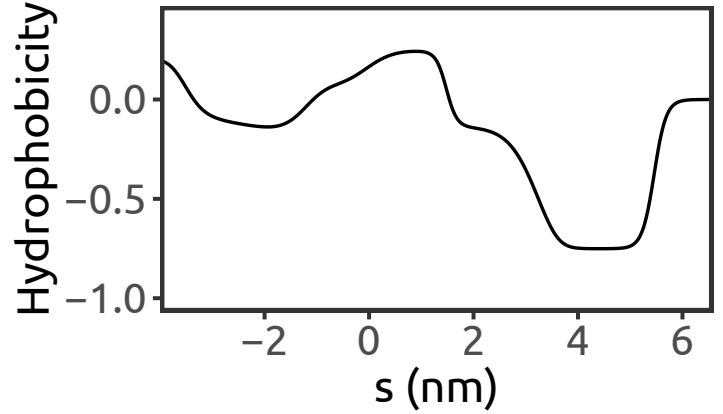
cryo-EM (4.22 Å)

Grieben et al., 2017

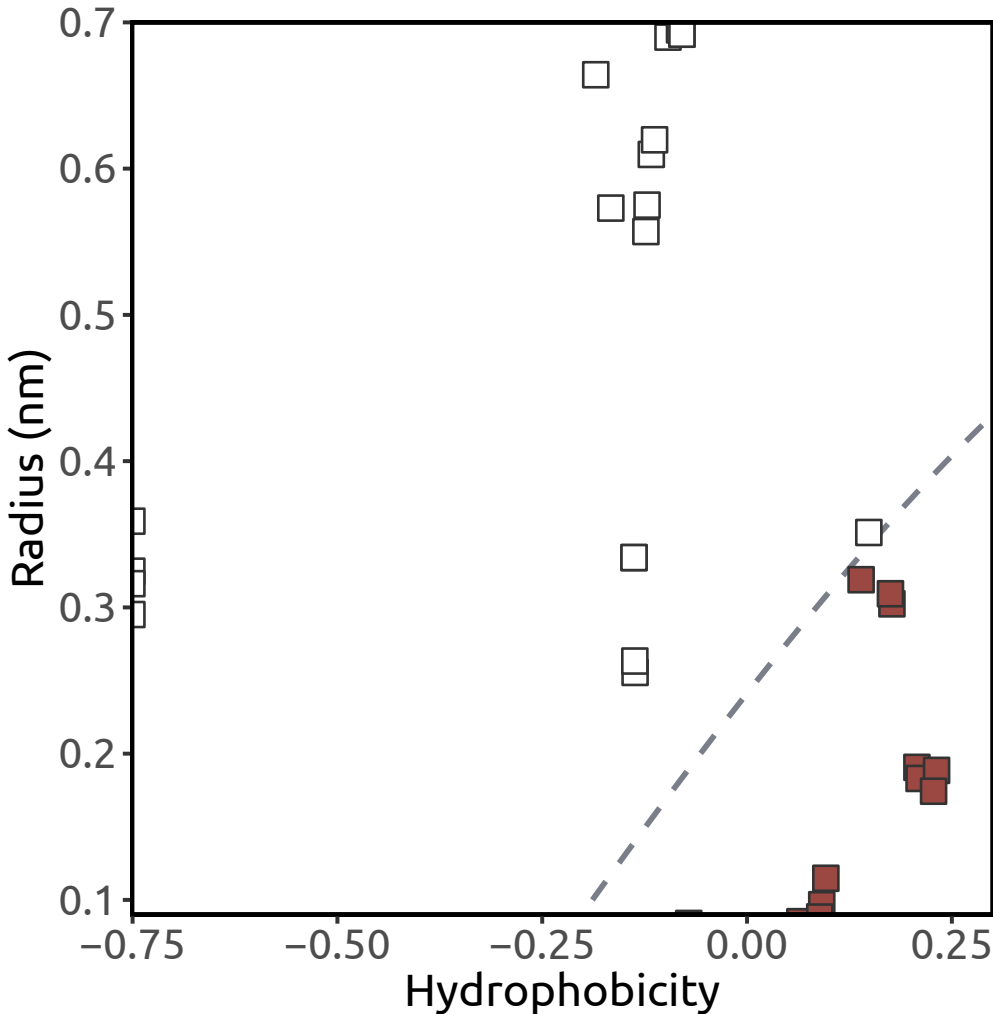
Pore radius



Hydrophobicity



Prediction



Heuristic score:

2.49 (n = 19)

Simulation result:

barrier to water

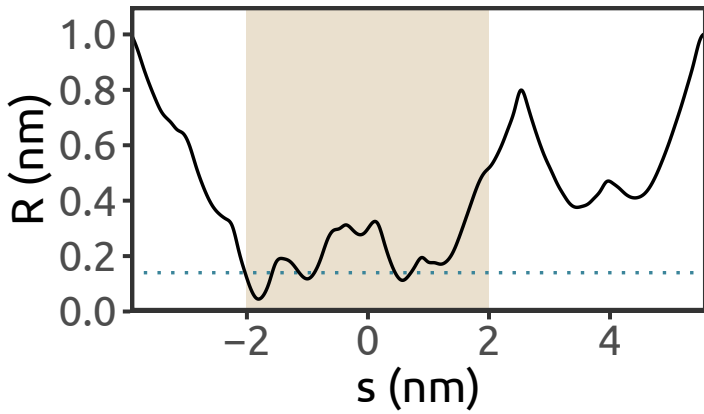
PKD2 (PDB ID: 5MKE)

Homo sapiens

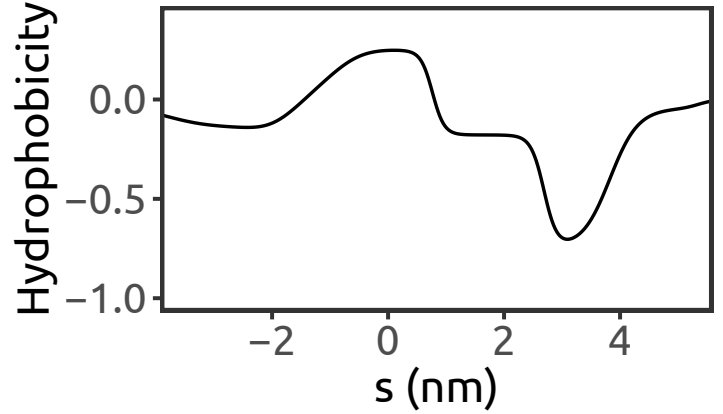
cryo-EM (4.3 Å)

Wilkes et al., 2017

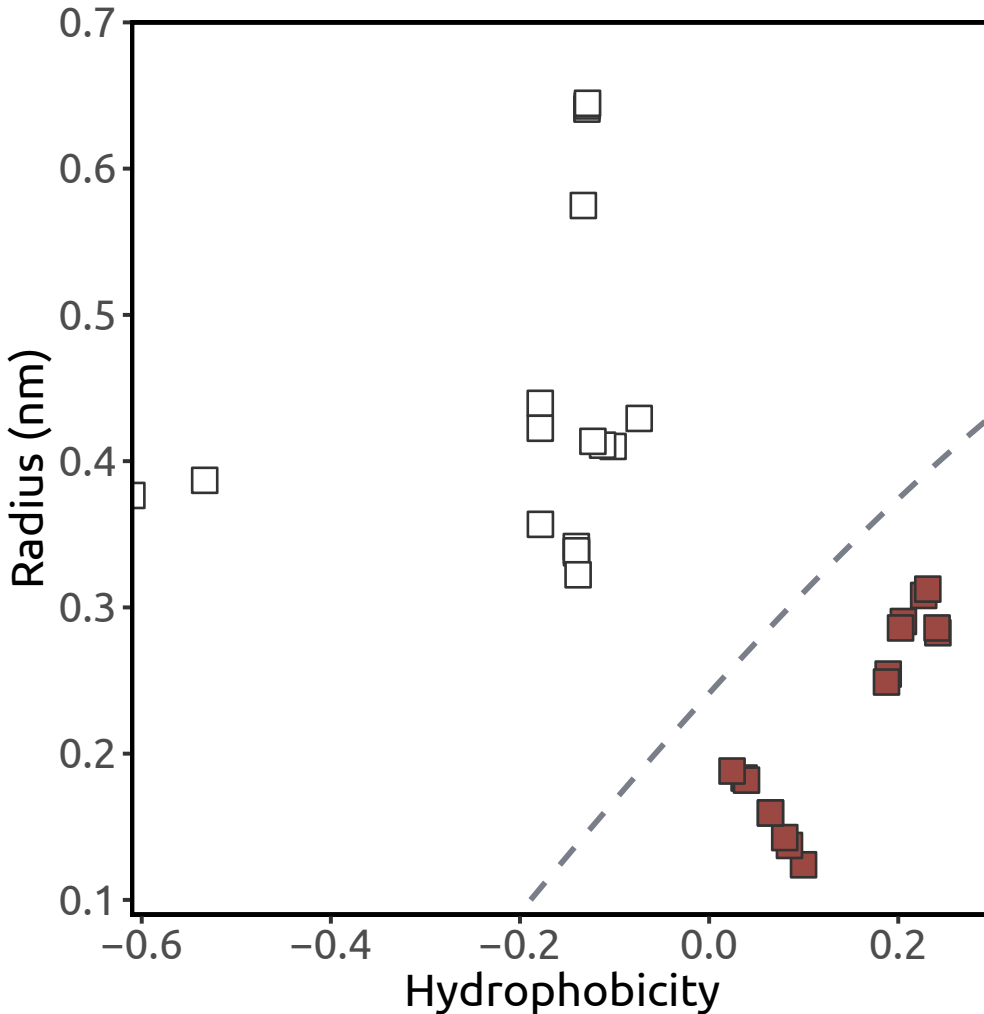
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.91 ($n = 20$)

Simulation result:

hydrated channel

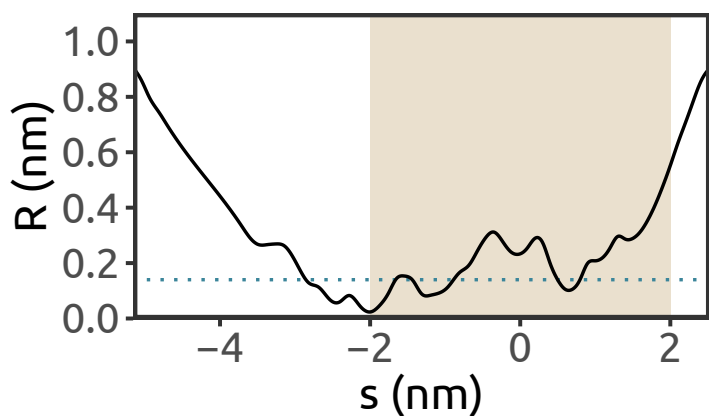
PKD2 (PDB ID: 5MKF)

Homo sapiens

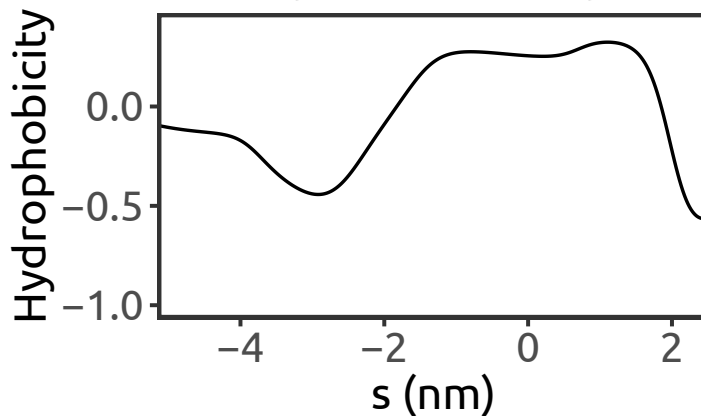
cryo-EM (4.2 Å)

Wilkes et al., 2017

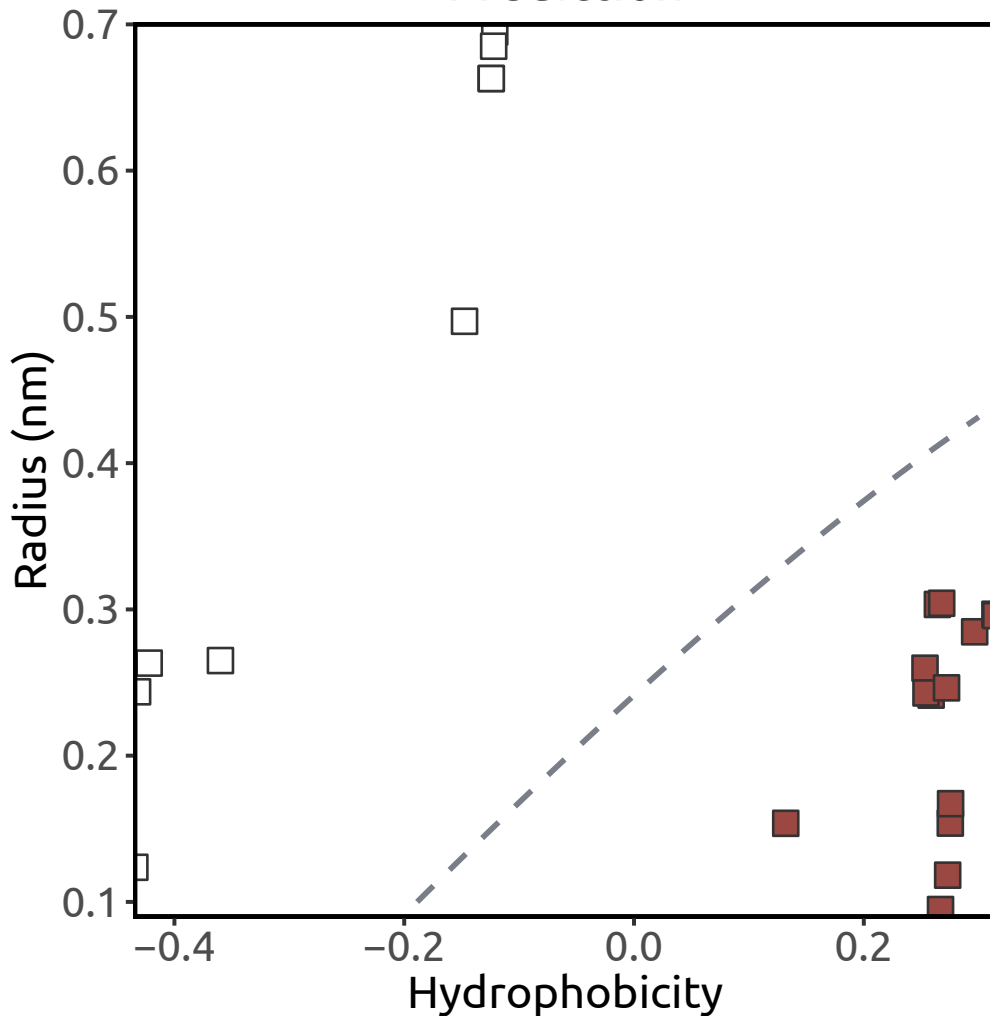
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.5 (n = 16)

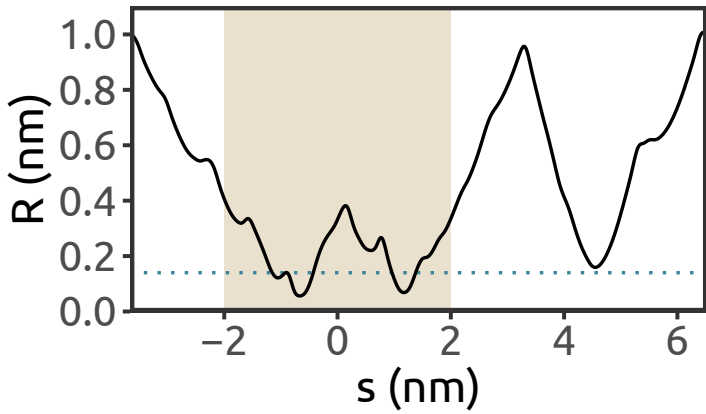
Simulation result:
barrier to water

PKD2 (PDB ID: 5T4D)

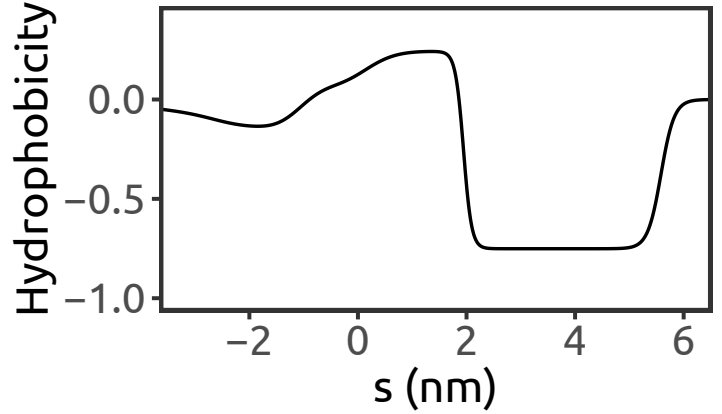
Homo sapiens
cryo-EM (3 Å)

Shen et al., 2016

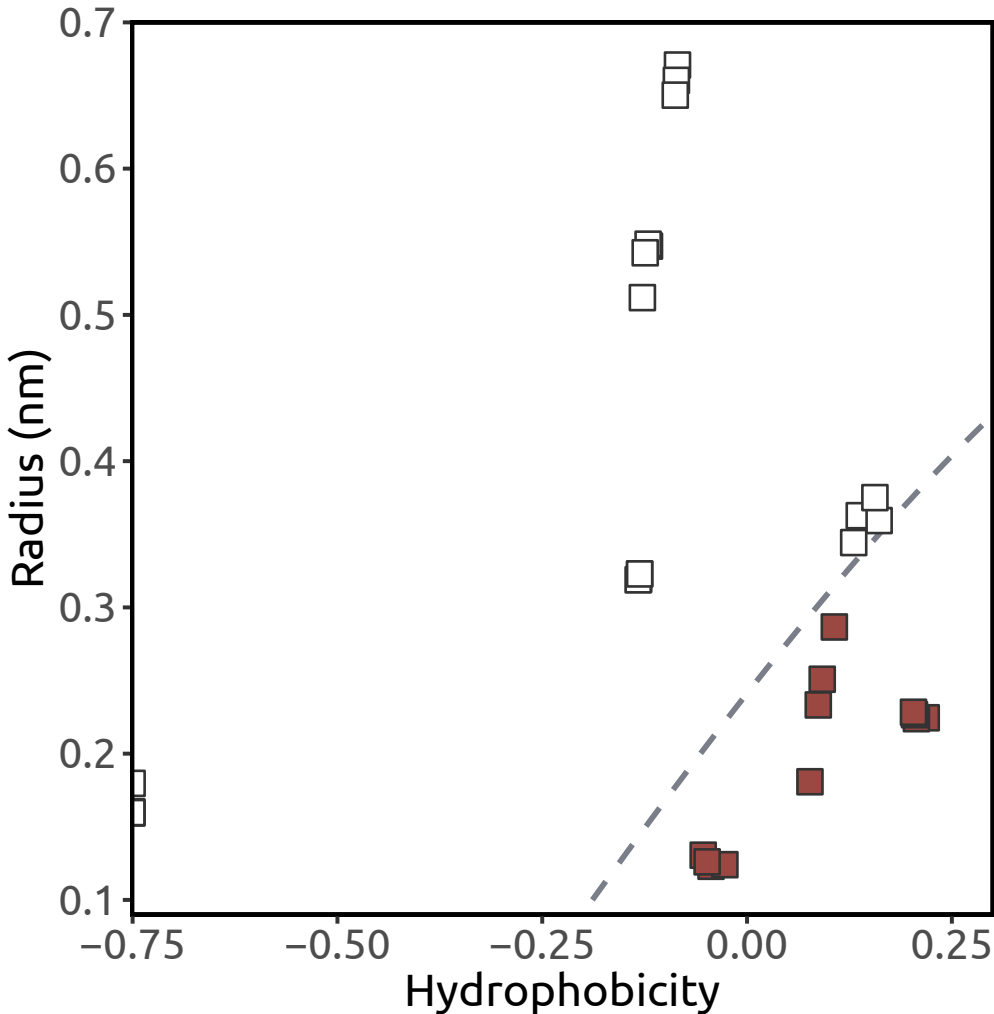
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.71 ($n = 16$)

Simulation result:
barrier to water

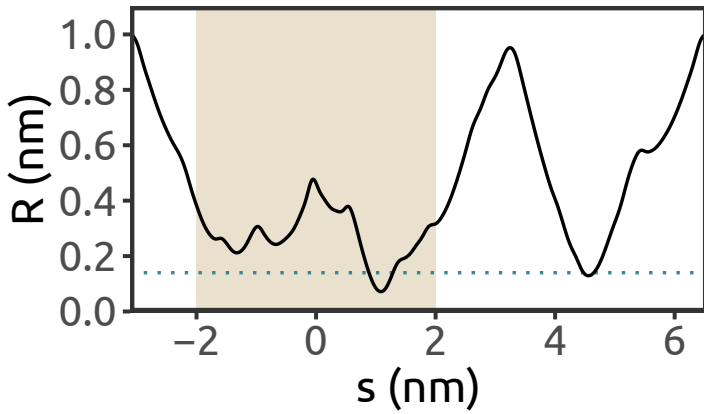
PKD2 (PDB ID: 6D1W)

Homo sapiens

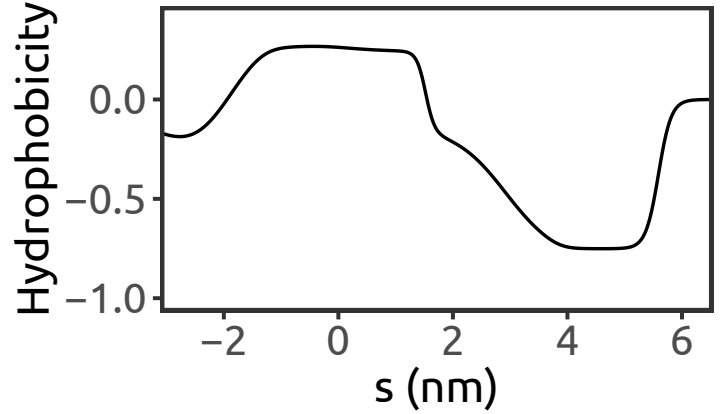
cryo-EM (3.54 Å)

Zheng et al., 2018

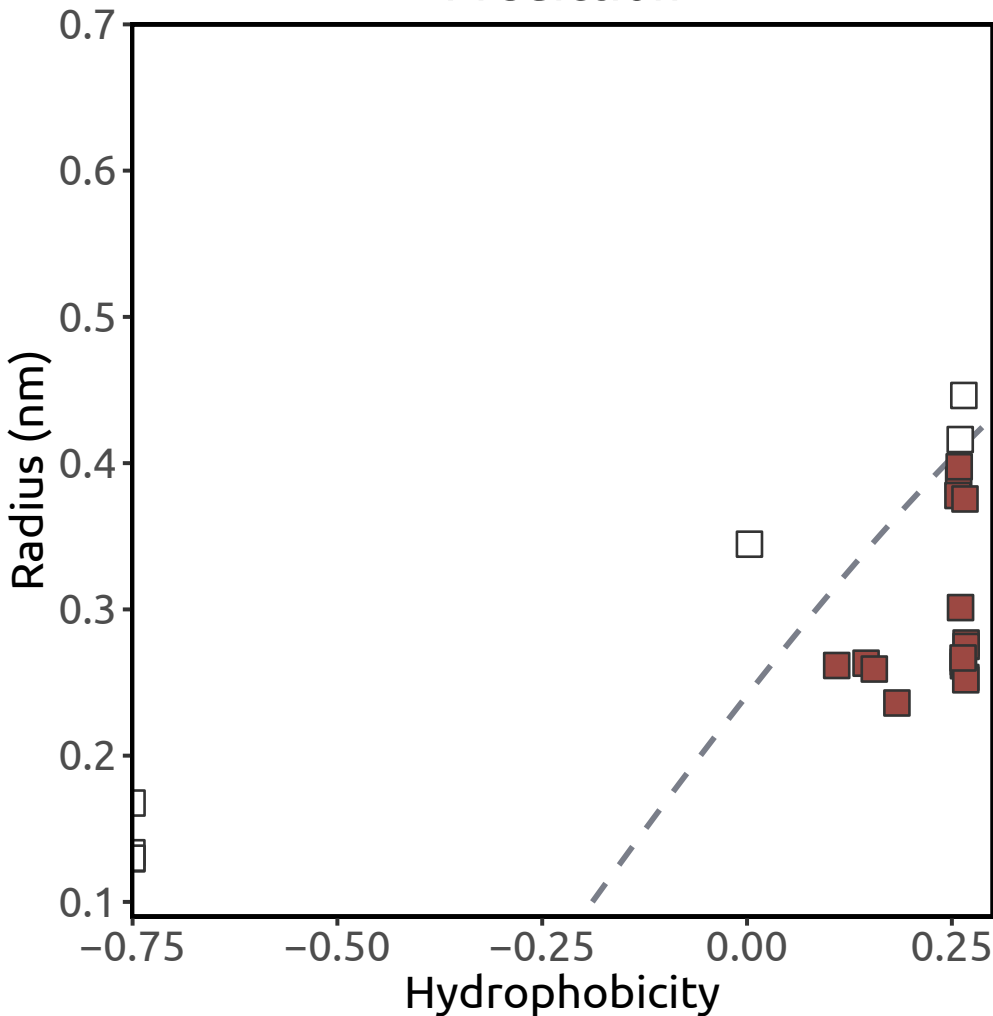
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.37 (n = 16)

Simulation result:

barrier to water

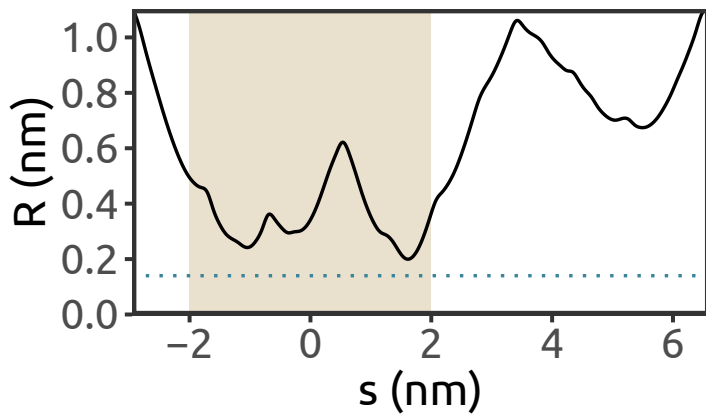
PKD2L1 (PDB ID: 5Z1W)

Mus musculus

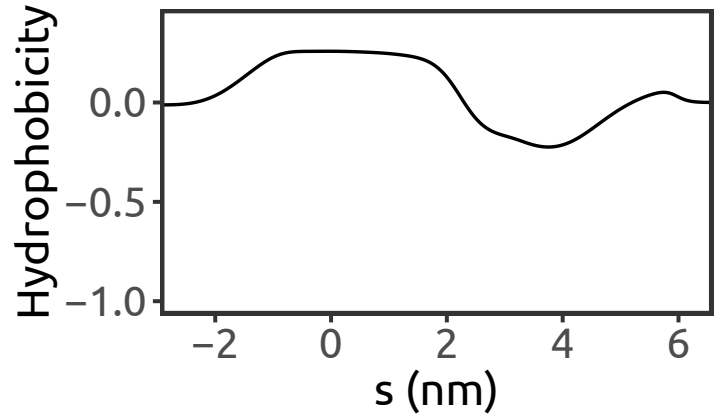
cryo-EM (3.38 Å)

Su et al., 2018

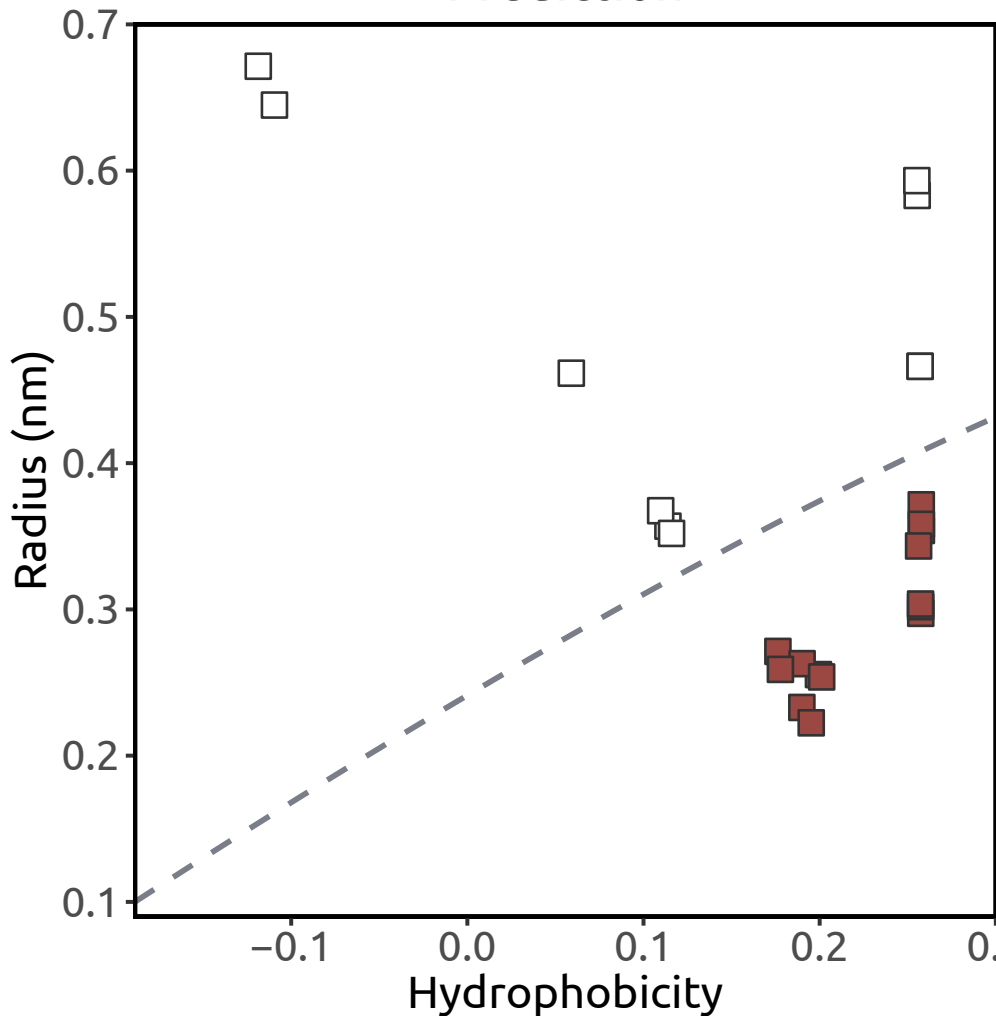
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.15 (n = 14)

Simulation result:

barrier to water

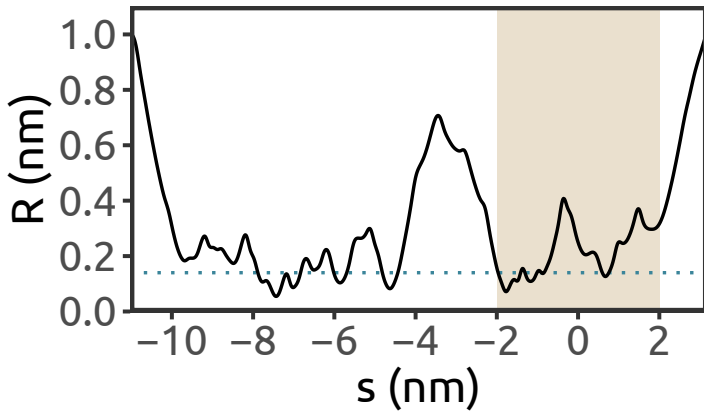
TRPA1 (PDB ID: 3J9P)

Homo sapiens

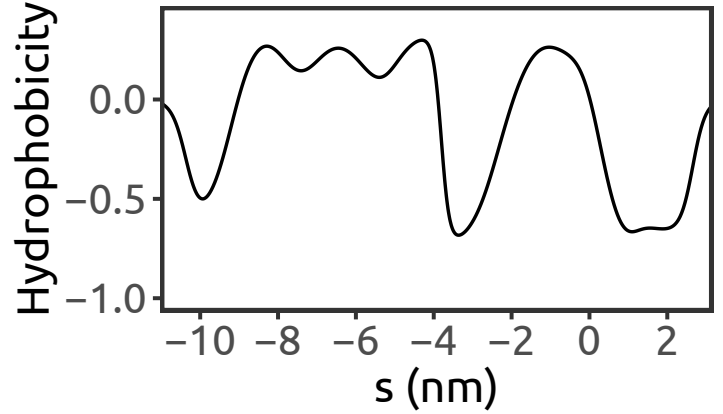
cryo-EM (4.24 Å)

Paulsen et al., 2015

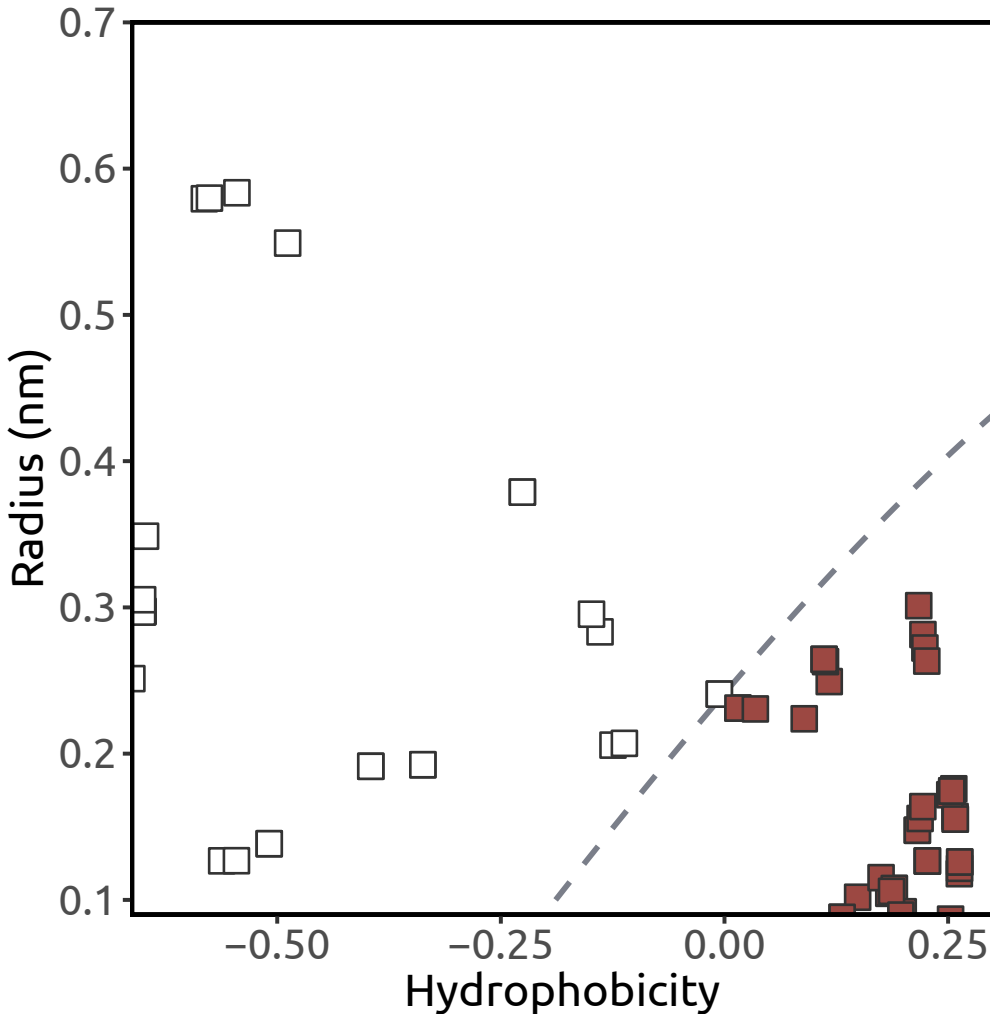
Pore radius



Hydrophobicity



Prediction



Heuristic score:

8.6 (n = 48)

Simulation result:

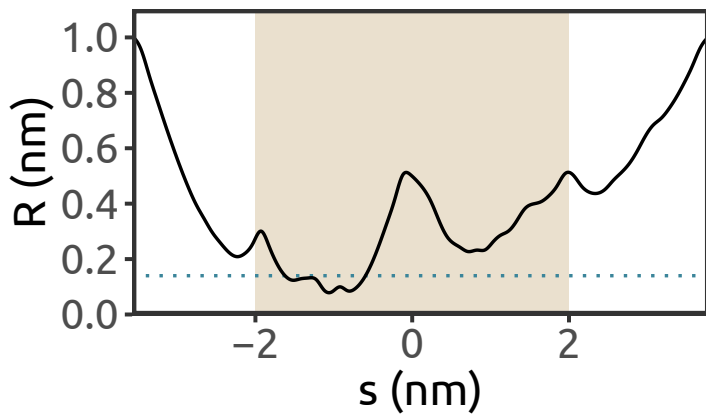
barrier to water

TRPM2 (PDB ID: 6CO7)

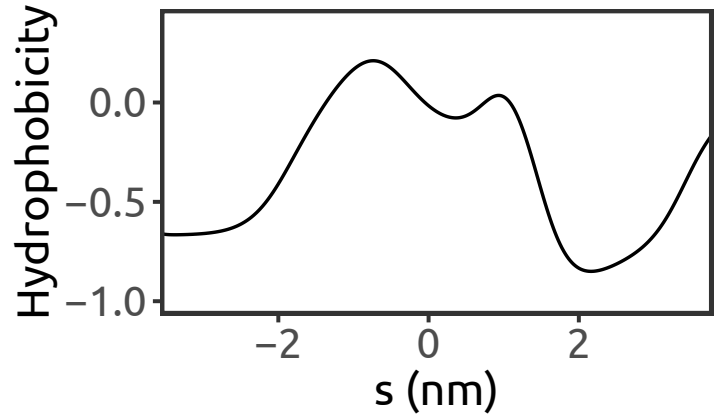
Nematostella vectensis
cryo-EM (3.07 Å)

Zhang et al., 2018

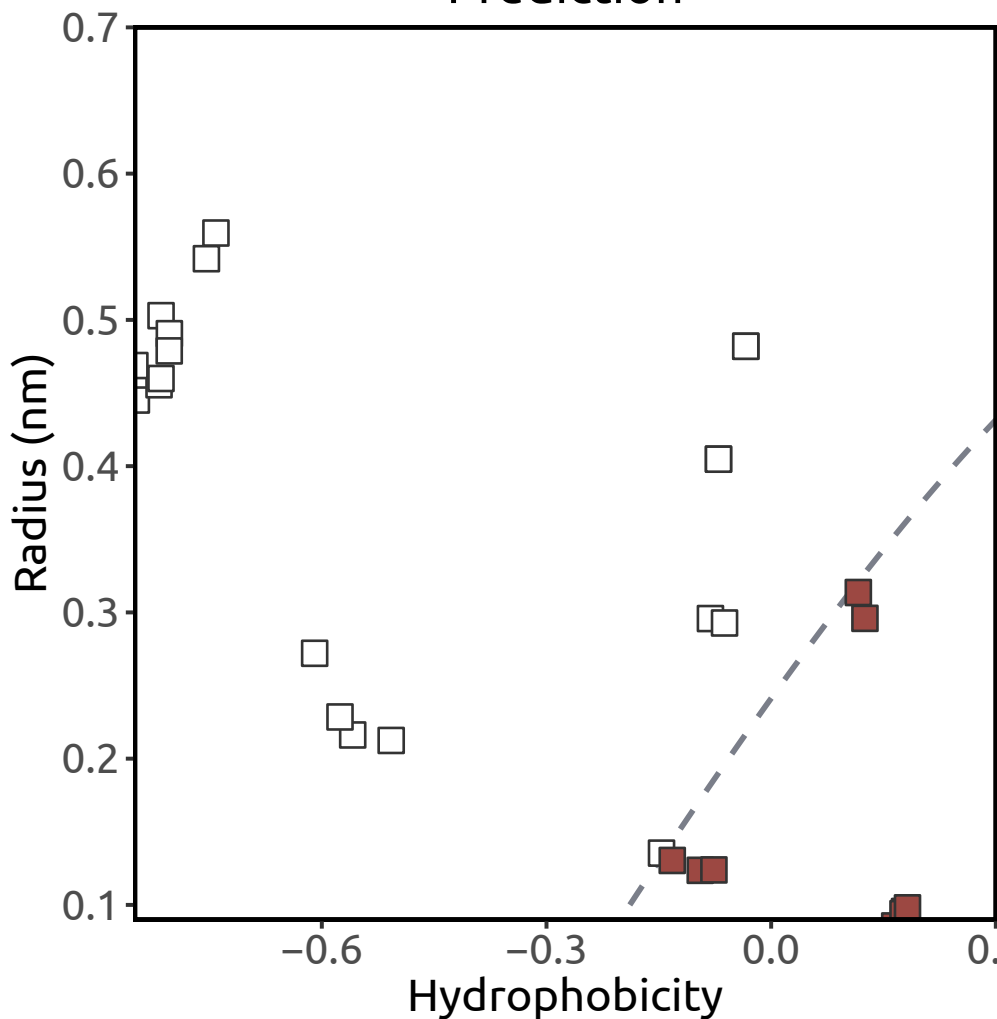
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.02 (n = 9)

Simulation result:
barrier to water

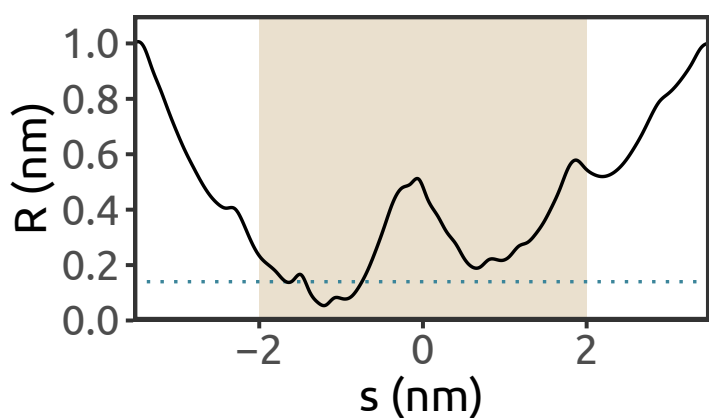
TRPM4 (PDB ID: 6BCL)

Mus musculus

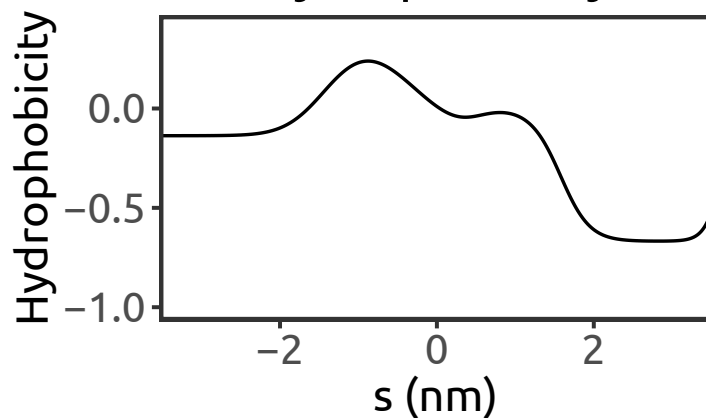
cryo-EM (3.54 Å)

Guo et al., 2017

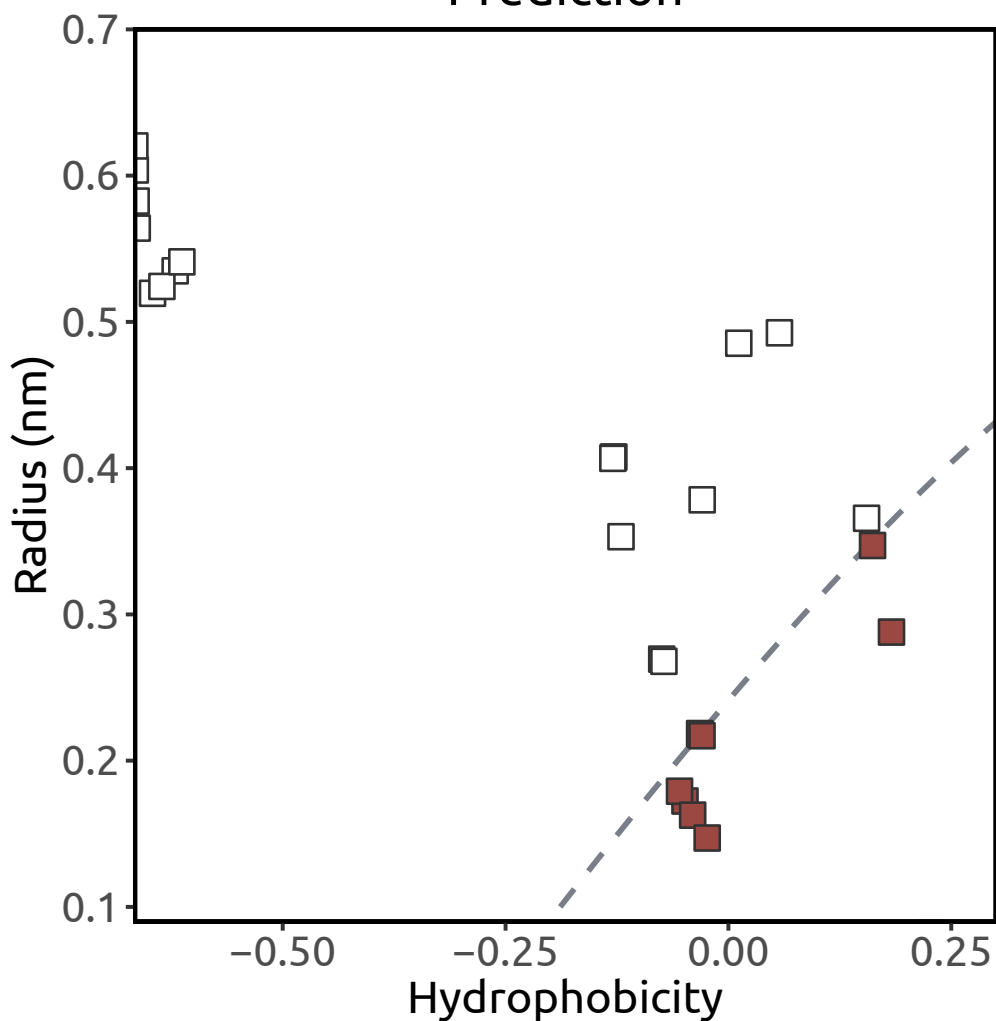
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.26 ($n = 13$)

Simulation result:
barrier to water

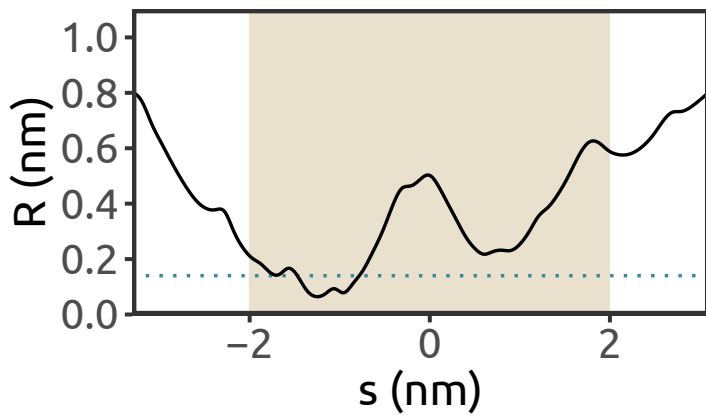
TRPM4 (PDB ID: 6BCQ)

Mus musculus

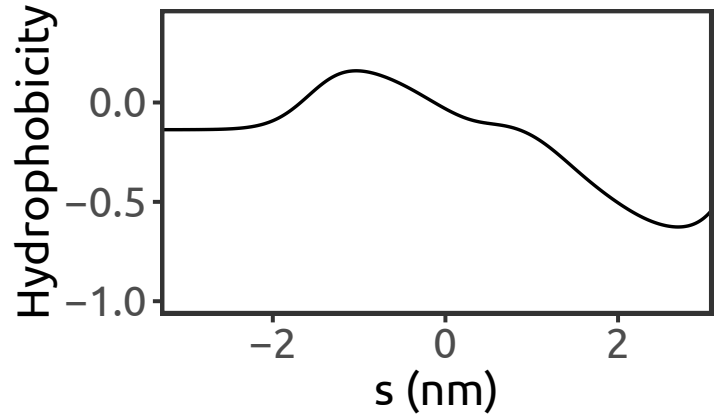
cryo-EM (3.25 Å)

Guo et al., 2017

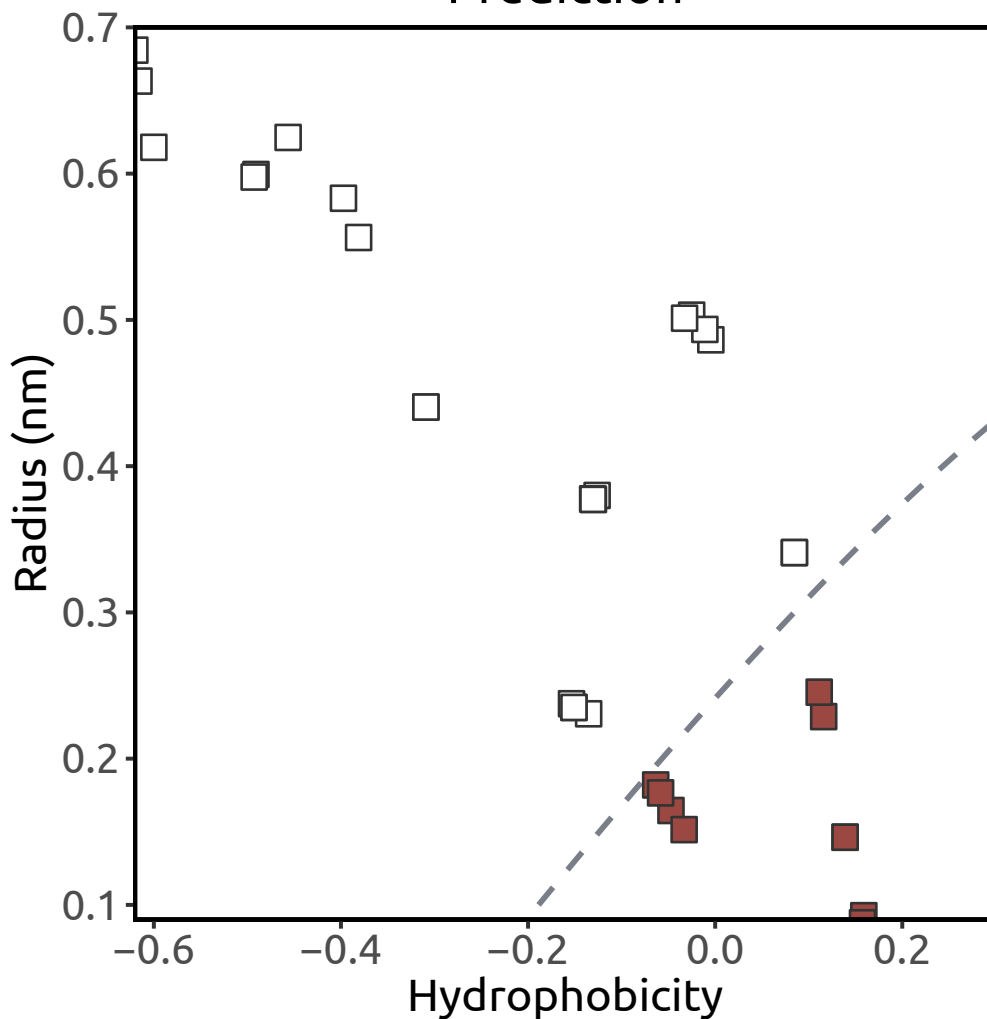
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.45 (n = 12)

Simulation result:
barrier to water

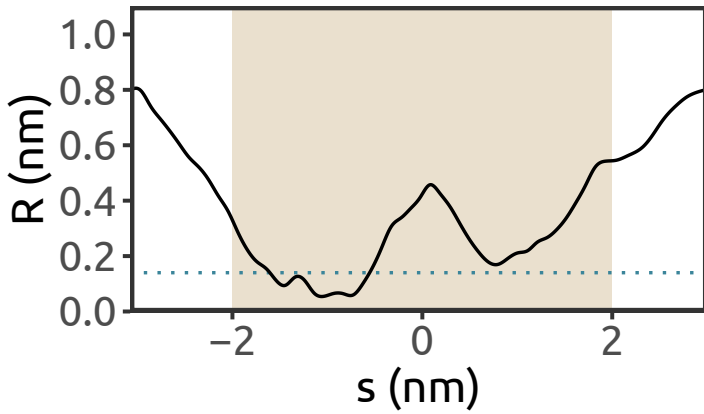
TRPM4 (PDB ID: 6BQR)

Homo sapiens

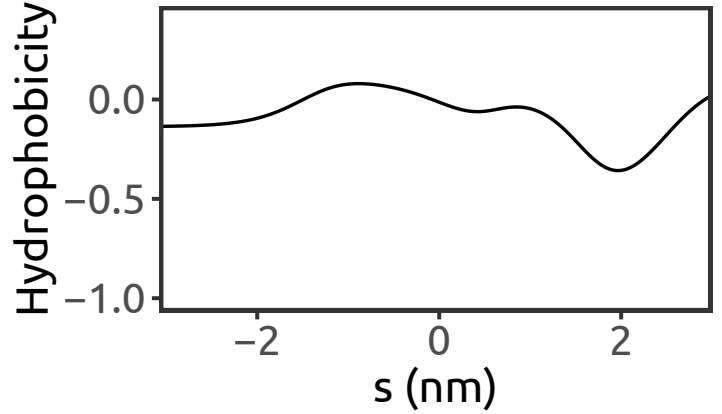
cryo-EM (3.2 Å)

Autzen et al., 2018

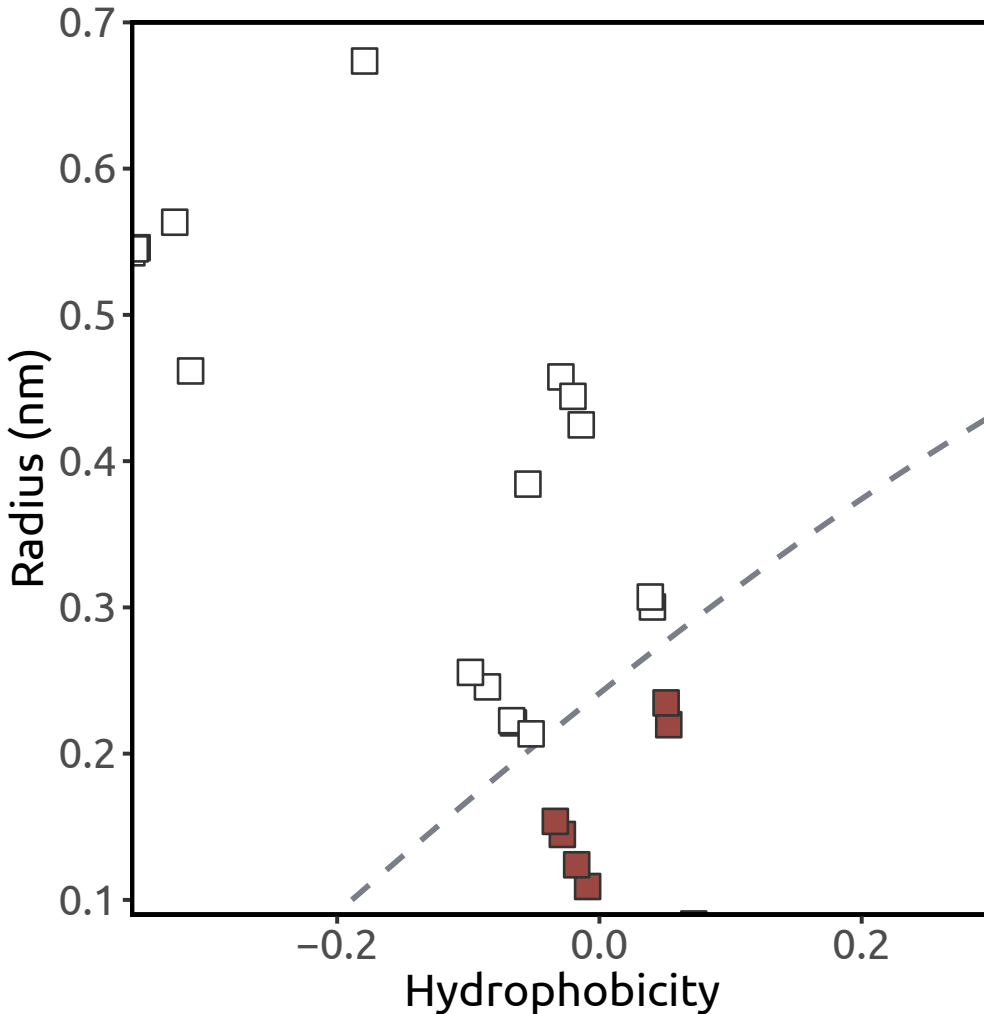
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.87 (n = 14)

Simulation result:

barrier to water

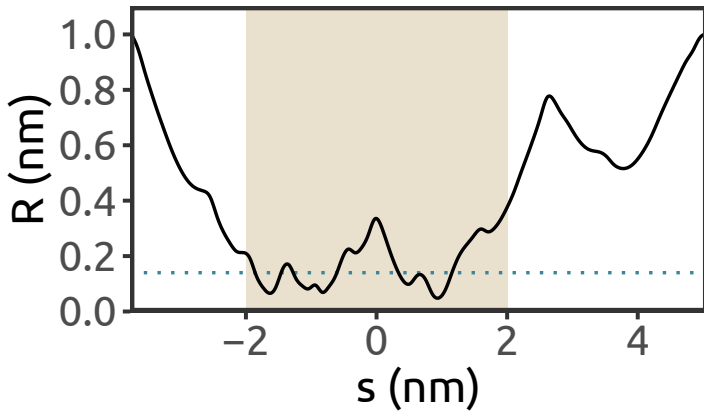
TRPML1 (PDB ID: 5WJ5)

Homo sapiens

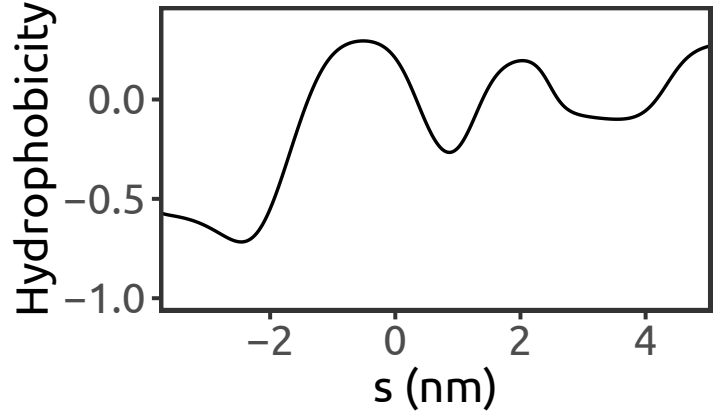
cryo-EM (3.72 Å)

Schmiege et al., 2017

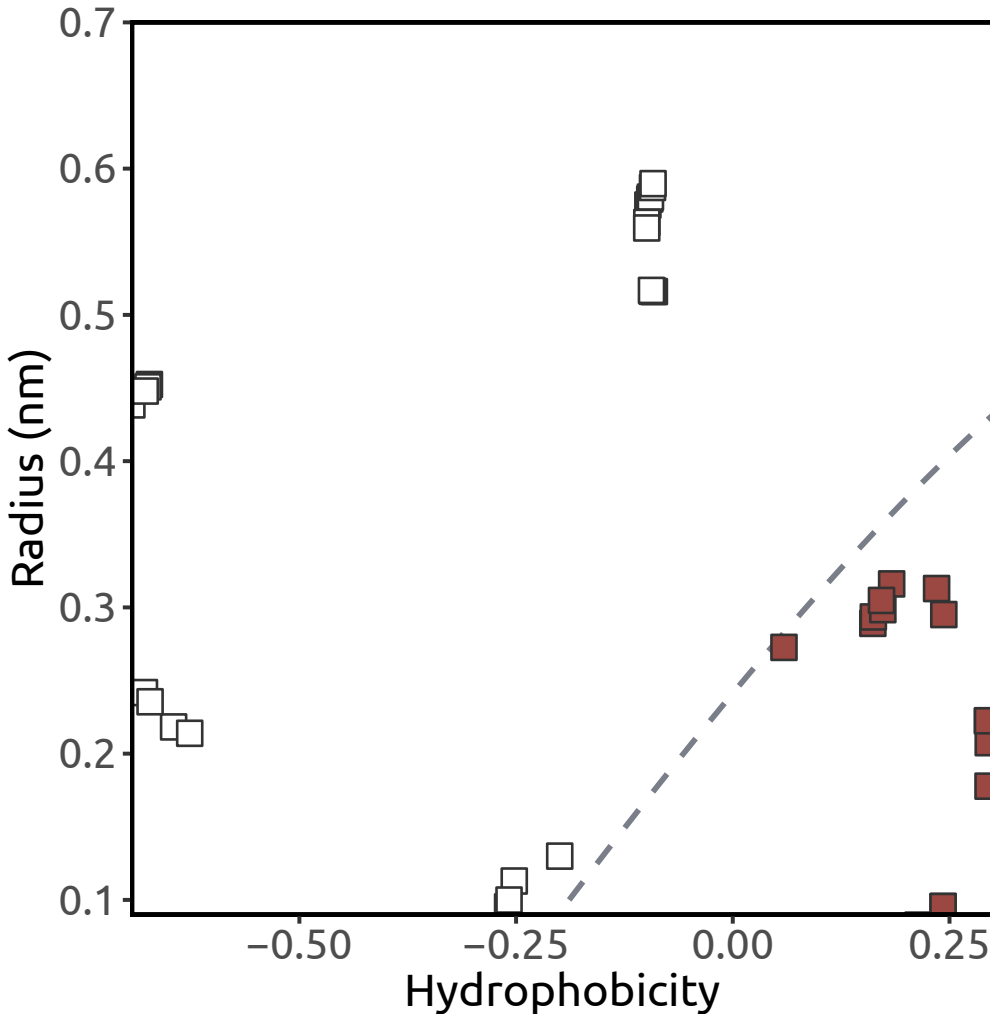
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.22 ($n = 17$)

Simulation result:
barrier to water

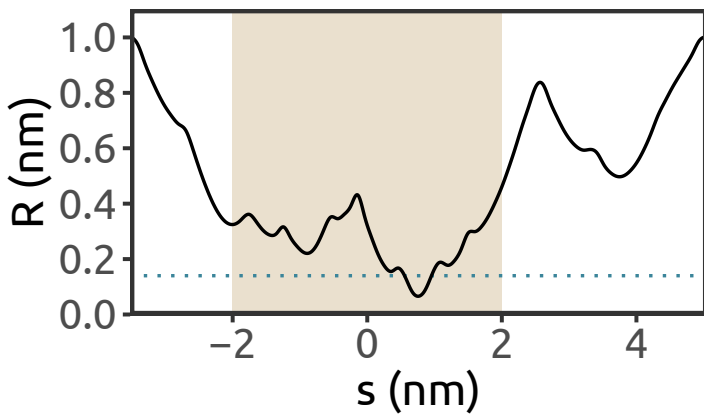
TRPML1 (PDB ID: 5WJ9)

Homo sapiens

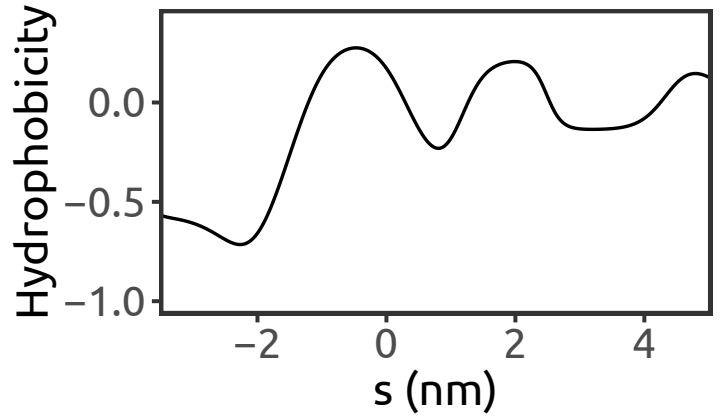
cryo-EM (3.49 Å)

Schmiege et al., 2017

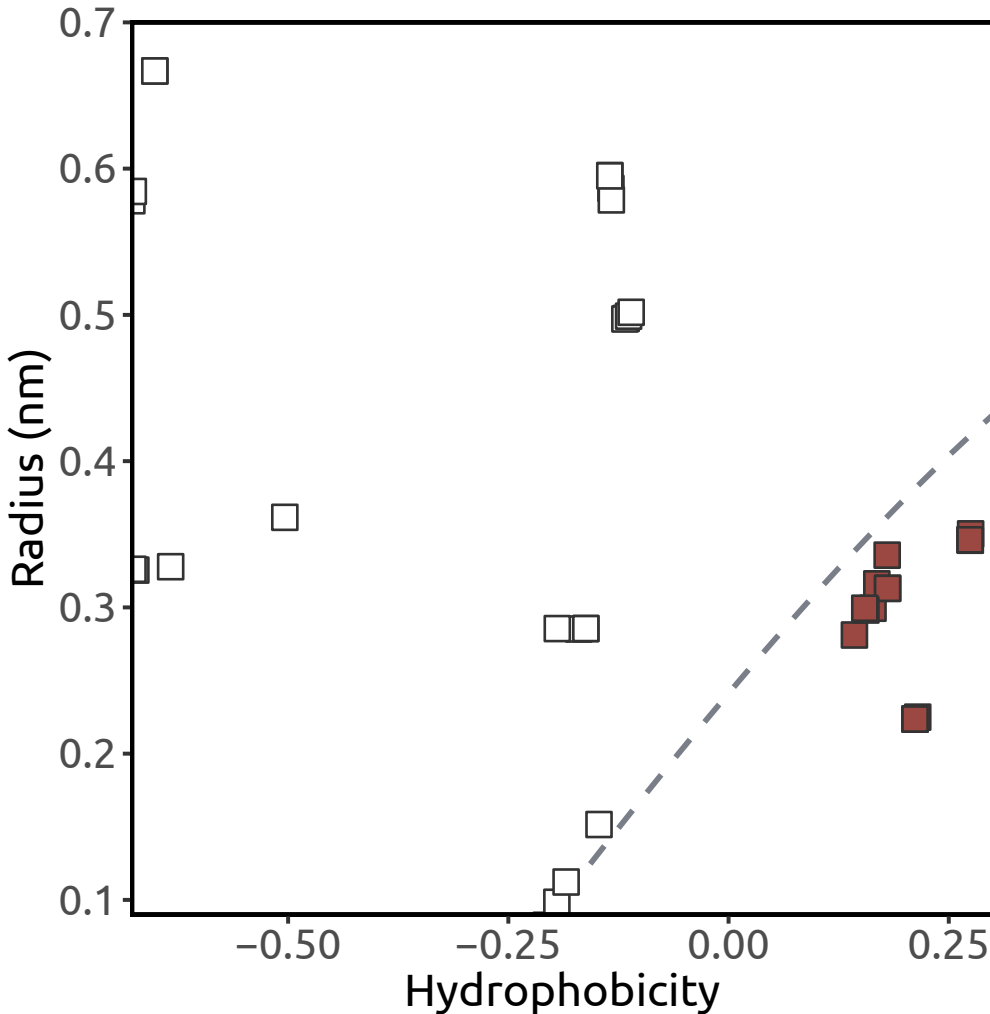
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.09 ($n = 16$)

Simulation result:

hydrated channel

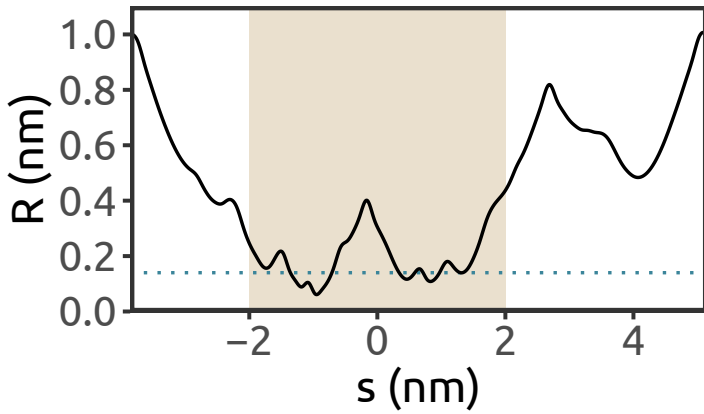
TRPML1 (PDB ID: 5WPV)

Mus musculus

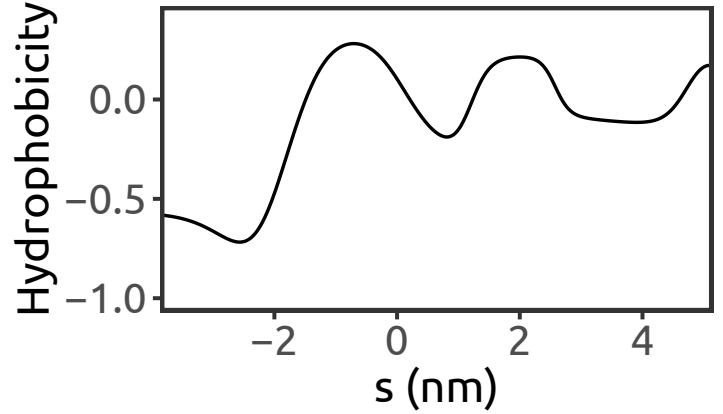
cryo-EM (3.59 Å)

Chen et al., 2017

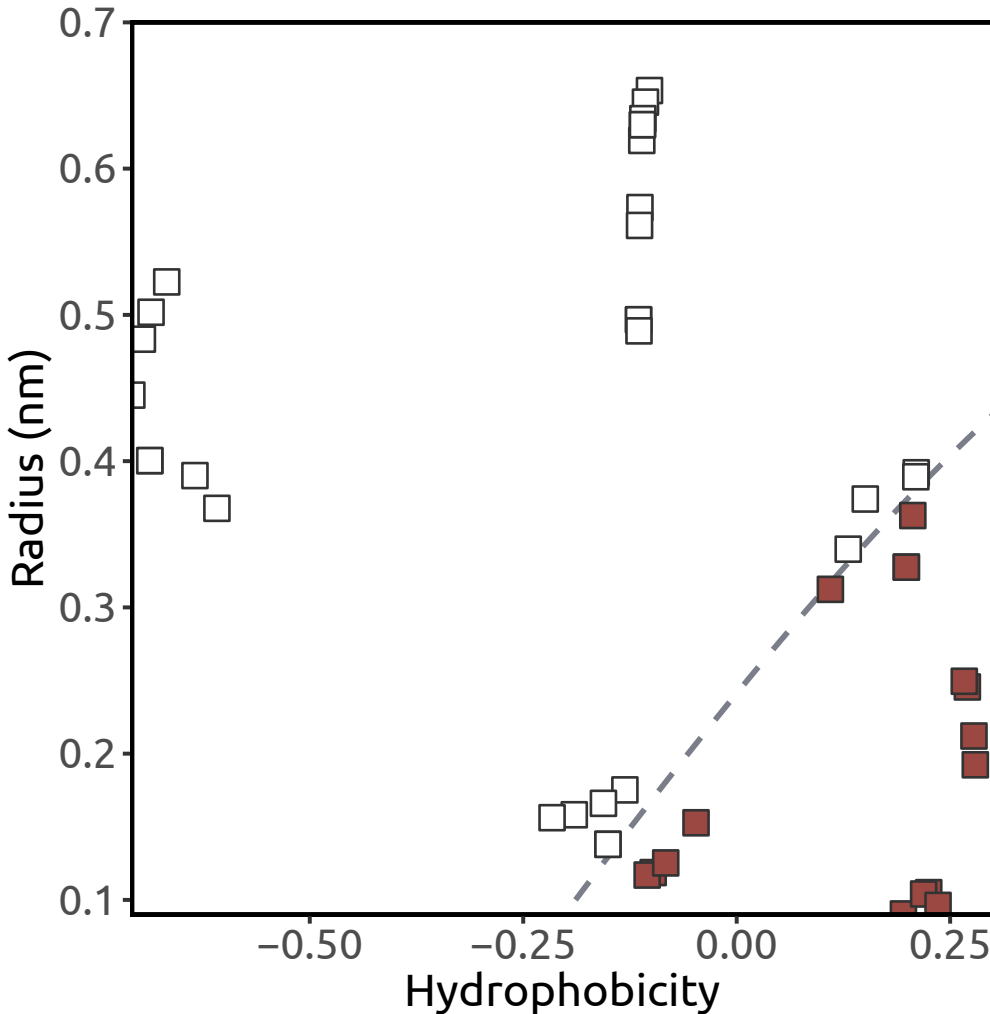
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.89 (n = 16)

Simulation result:

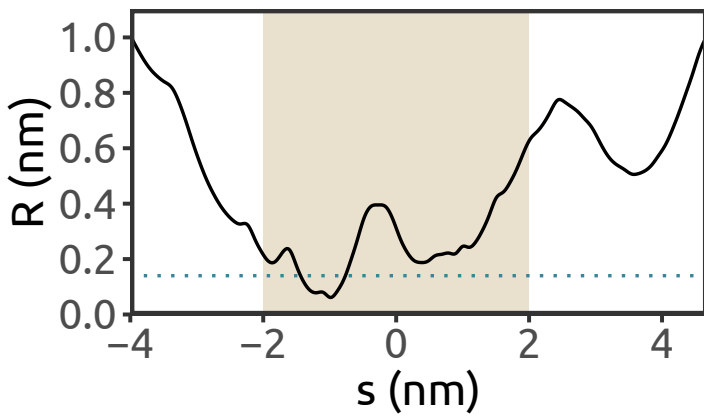
barrier to water

TRPML3 (PDB ID: 5W3S)

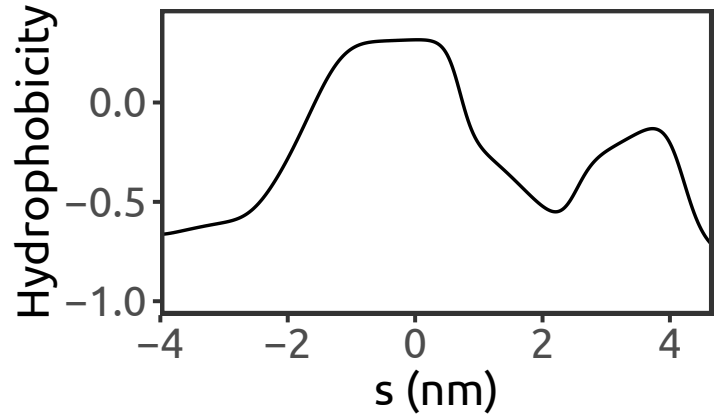
Callithrix jacchus
cryo-EM (2.94 Å)

Hirschi et al., 2017

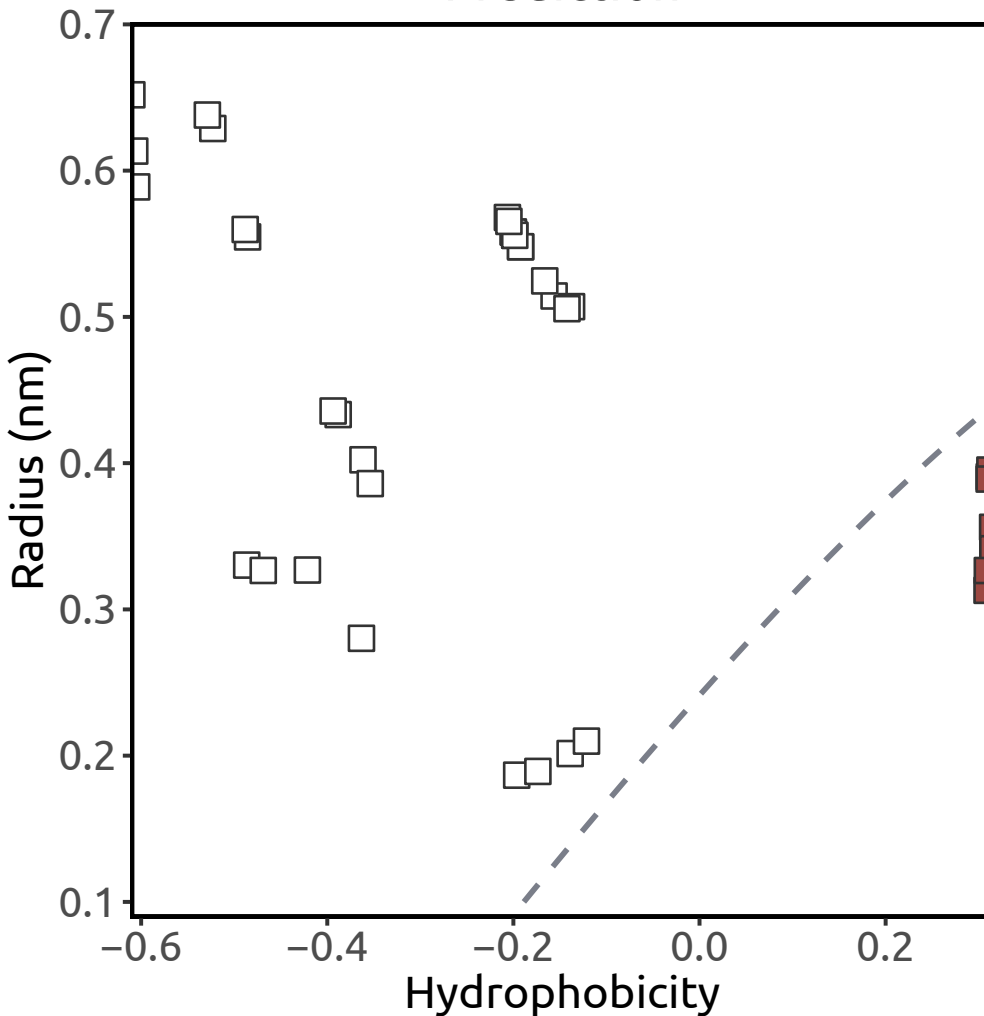
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.56 (n = 12)

Simulation result:
barrier to water

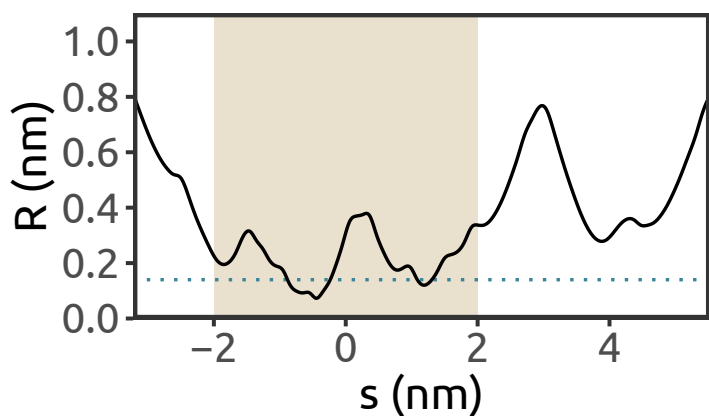
TRPML3 (PDB ID: 6AYE)

Homo sapiens

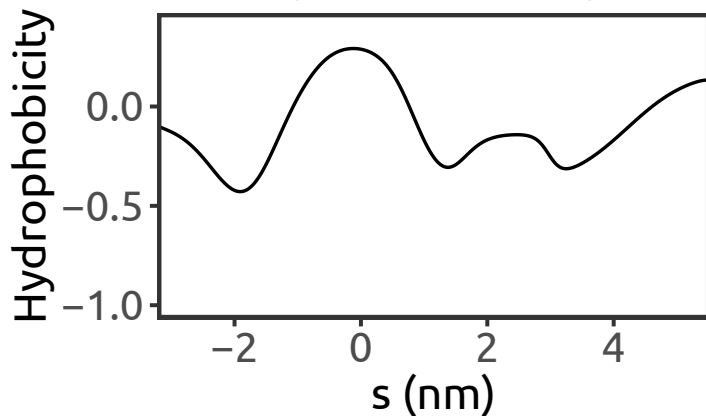
cryo-EM (4.06 Å)

Zhou et al., 2017

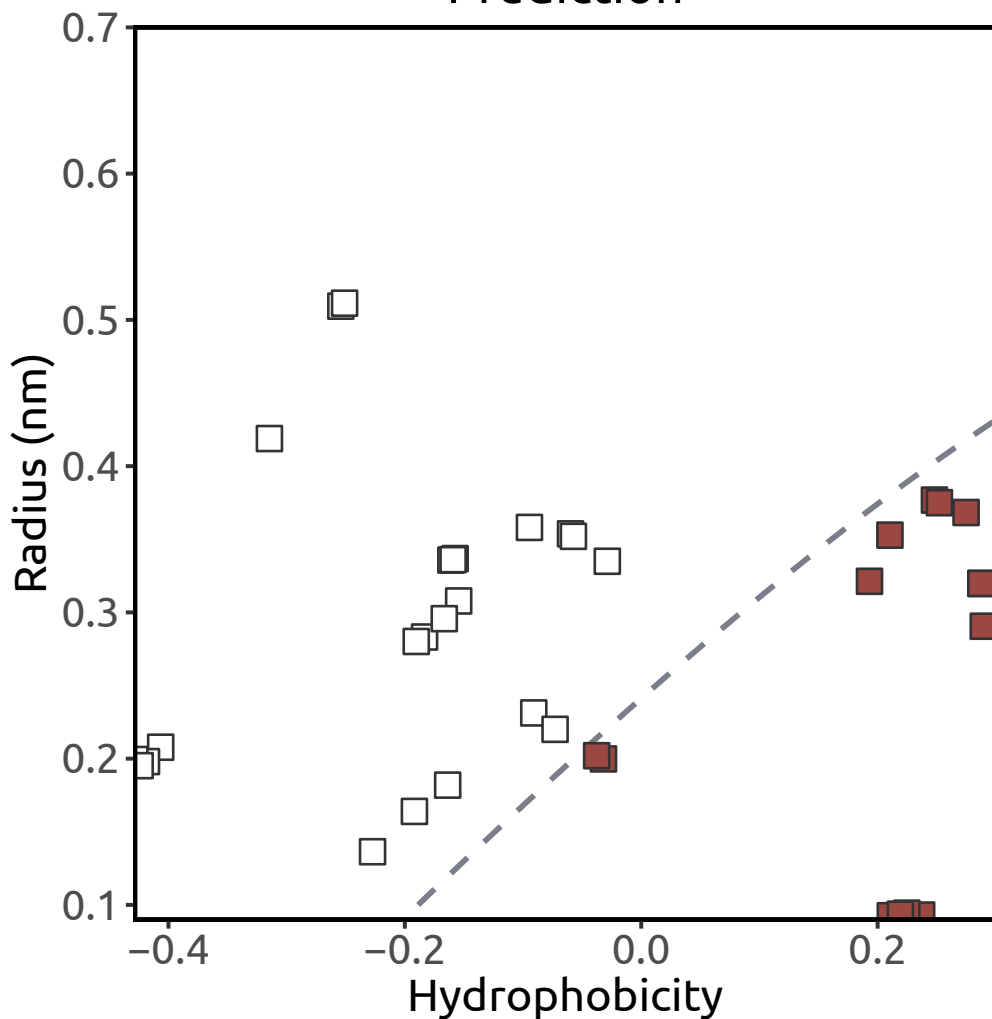
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.48 (n = 14)

Simulation result:
barrier to water

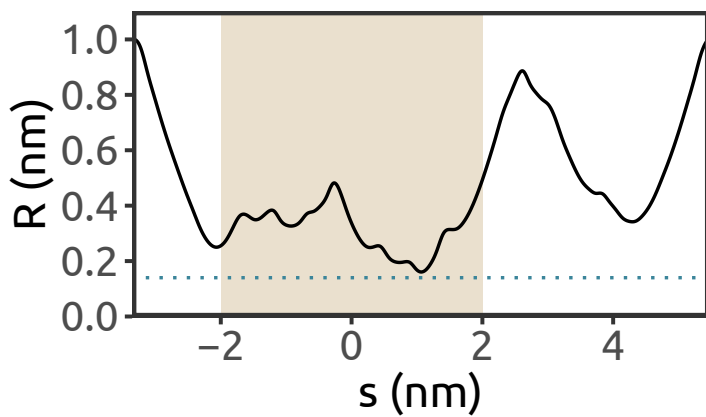
TRPML3 (PDB ID: 6AYF)

Homo sapiens

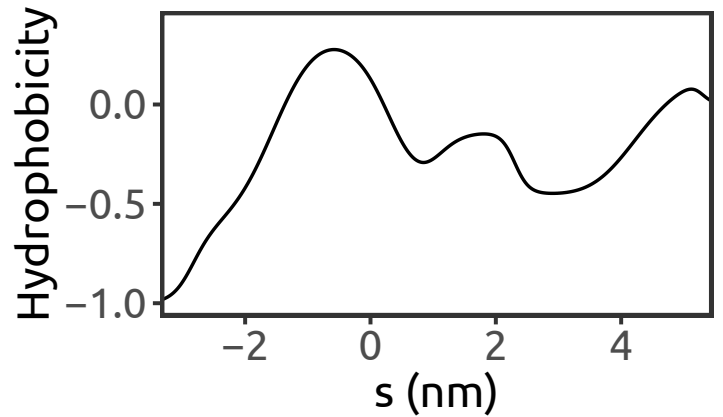
cryo-EM (3.62 Å)

Zhou et al., 2017

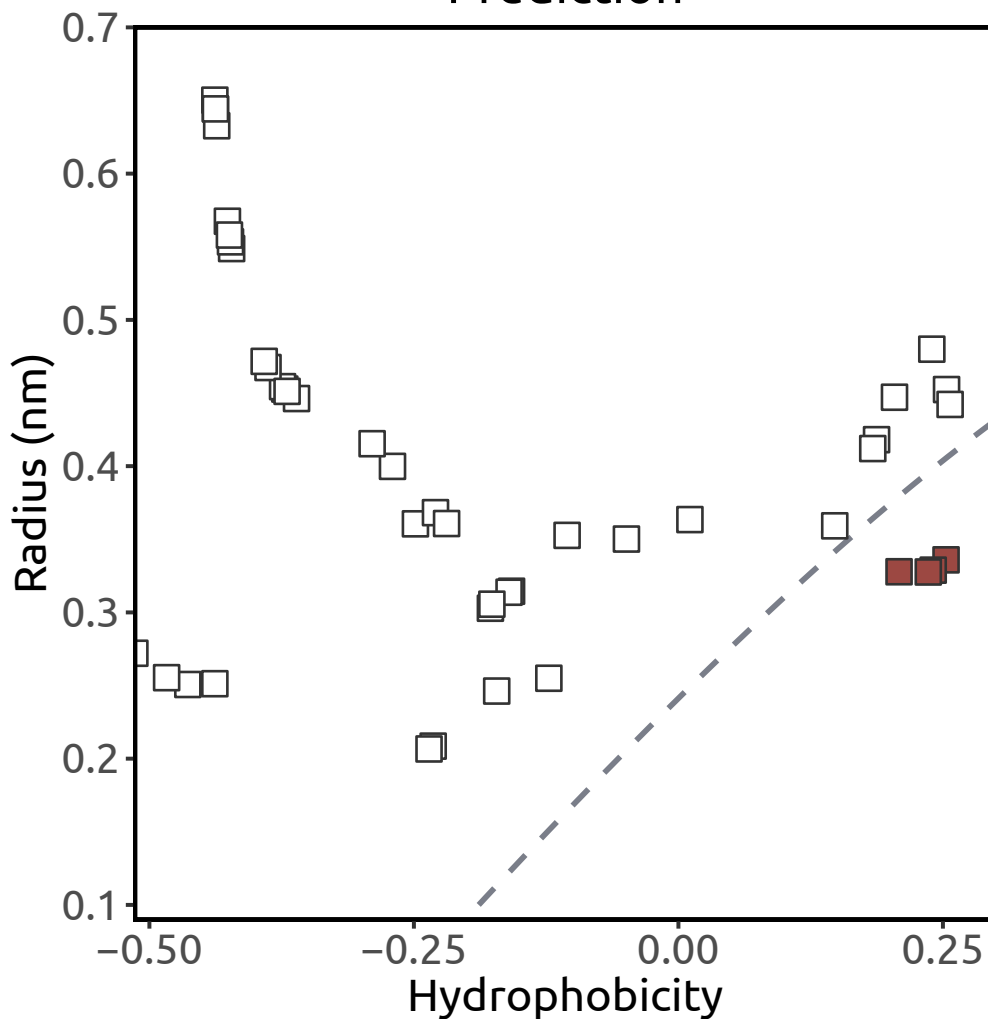
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.22 (n = 4)

Simulation result:
hydrated channel

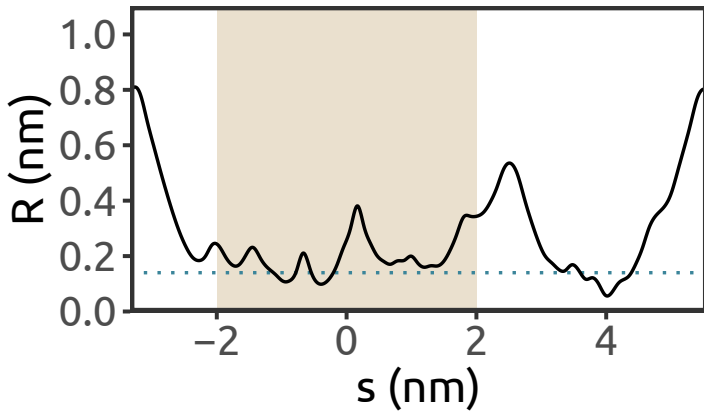
TRPML3 (PDB ID: 6AYG)

Homo sapiens

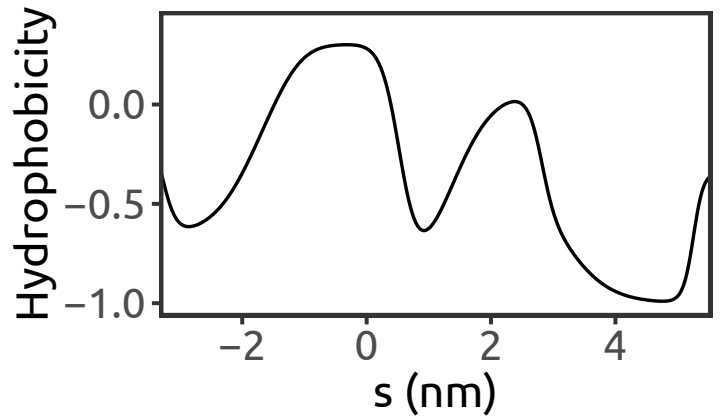
cryo-EM (4.65 Å)

Zhou et al., 2017

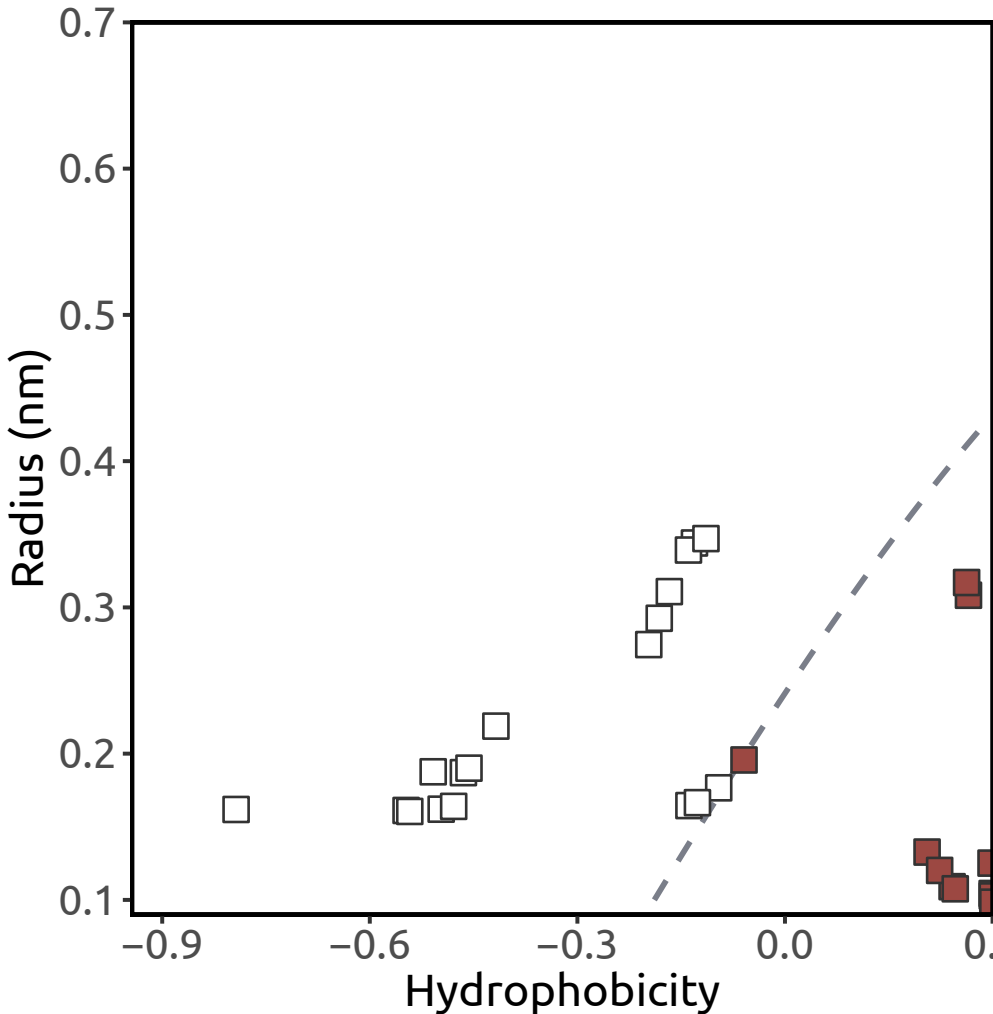
Pore radius



Hydrophobicity



Prediction



Heuristic score:

2.22 (n = 11)

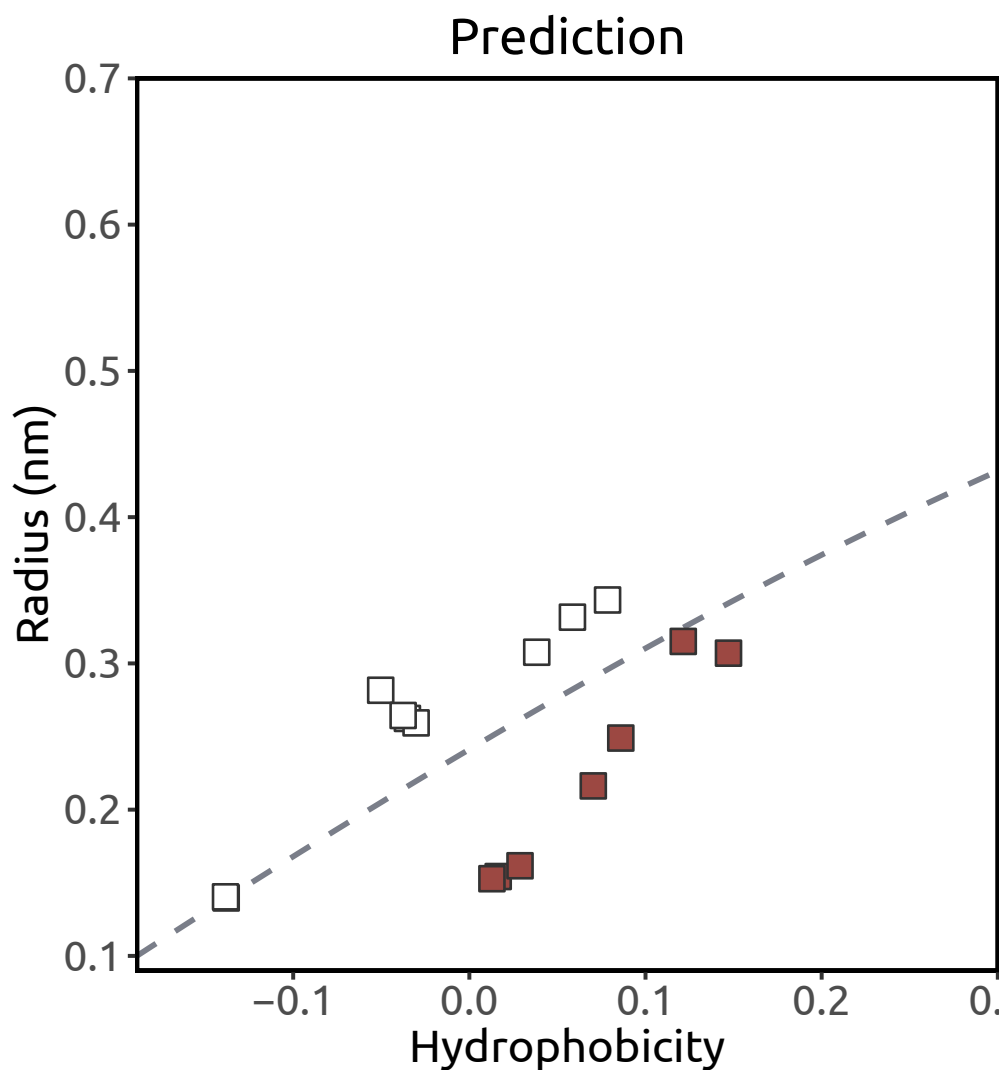
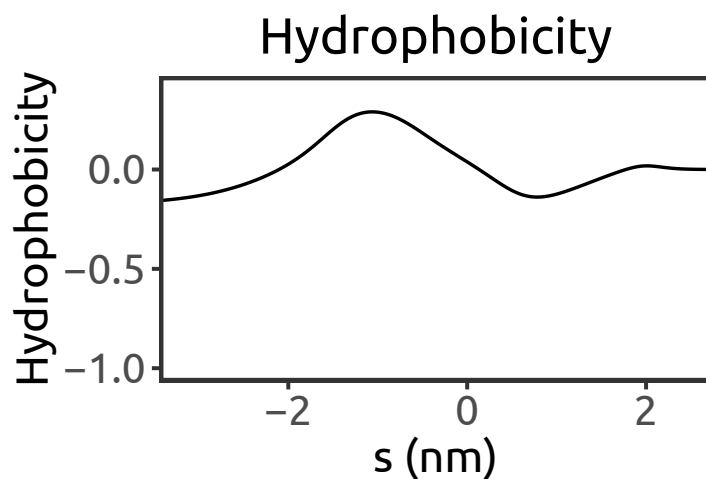
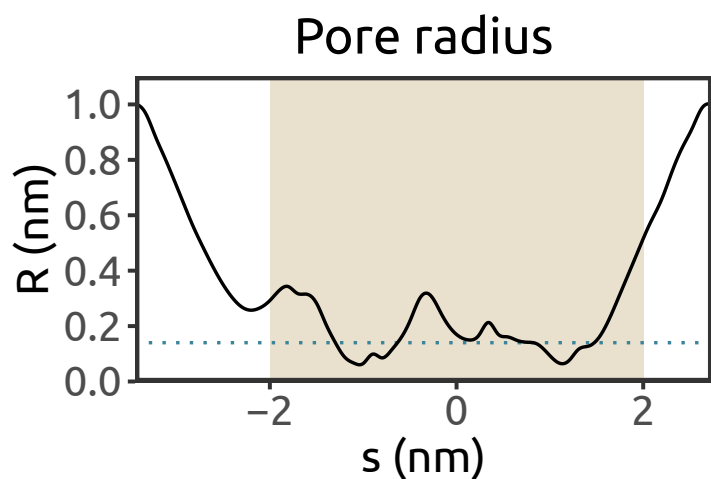
Simulation result:

barrier to water

TRPV1 (PDB ID: 3J5P)

Rattus norvegicus
cryo-EM (3.28 Å)

Liao et al., 2013



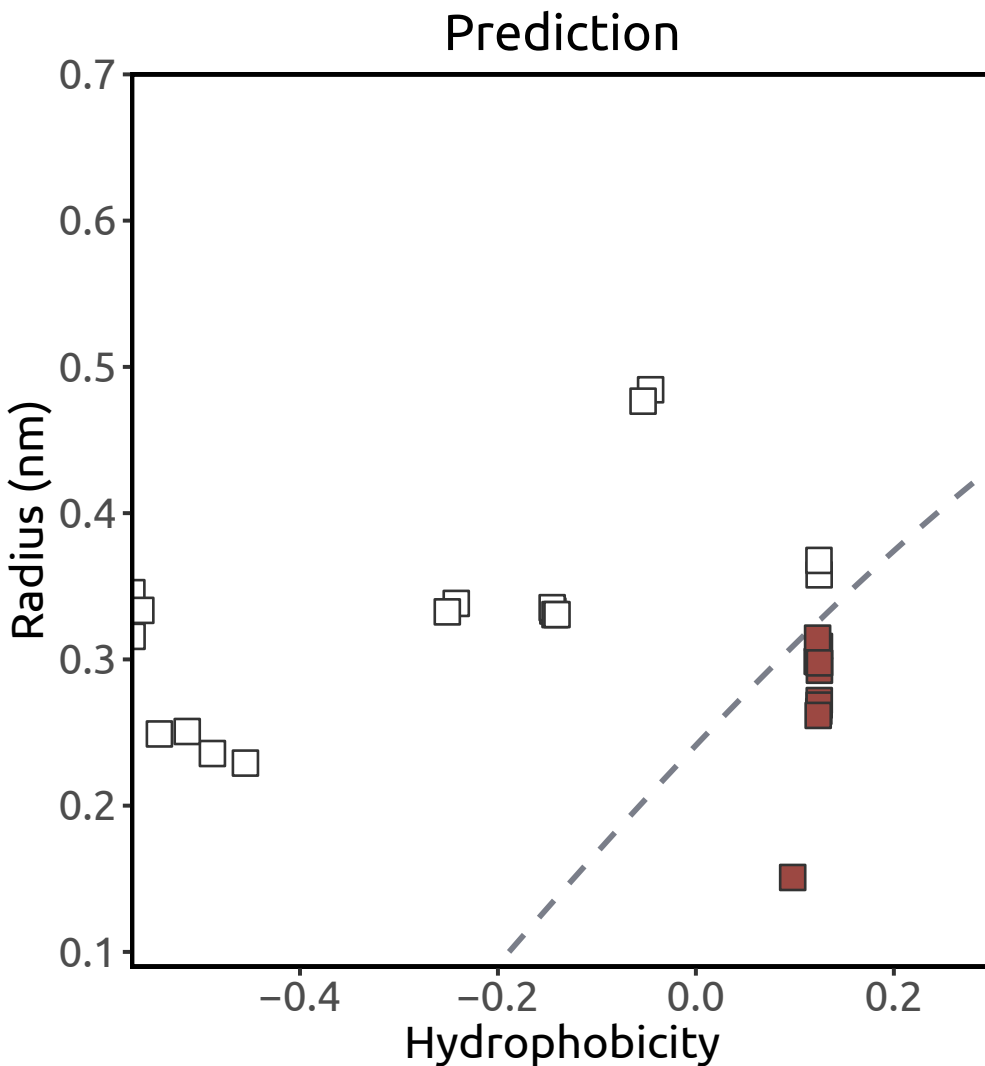
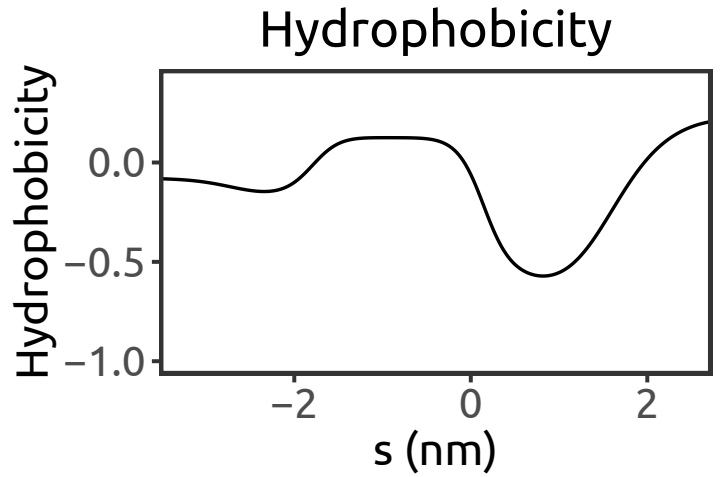
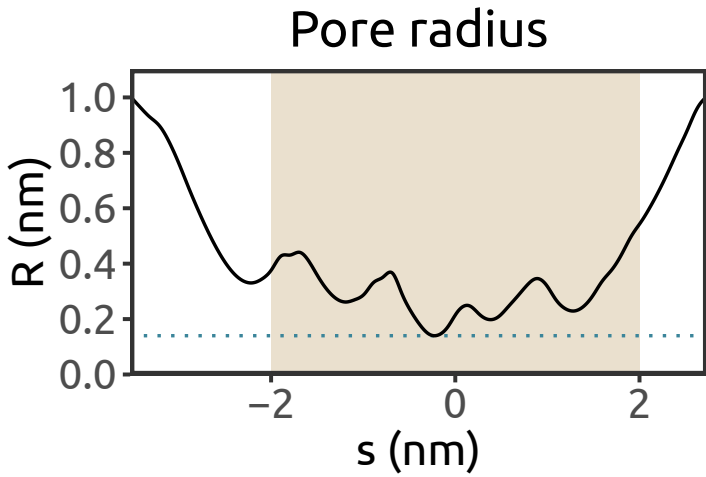
Heuristic score:
1.74 (n = 13)

Simulation result:
barrier to water

TRPV1 (PDB ID: 3J5Q)

Rattus norvegicus
cryo-EM (3.8 Å)

Cao et al., 2013



Heuristic score:
0.45 (n = 11)

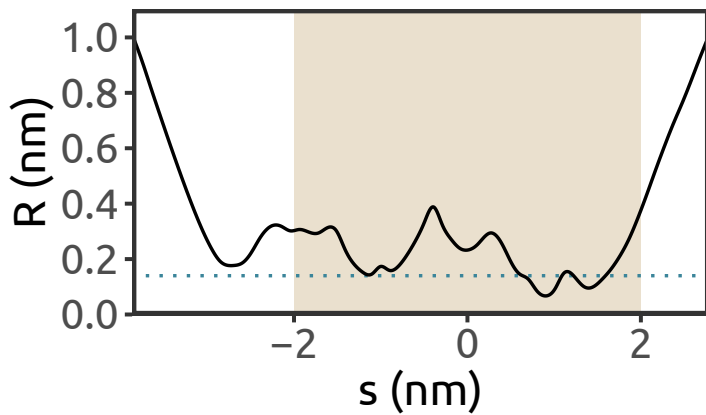
Simulation result:
barrier to water

TRPV1 (PDB ID: 3J5R)

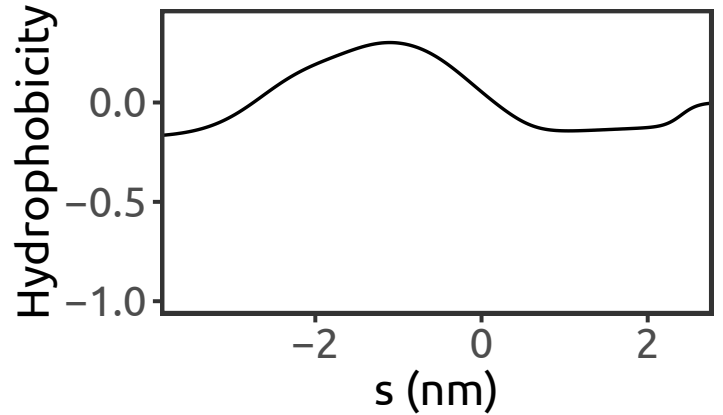
Rattus norvegicus
cryo-EM (4.2 Å)

Cao et al., 2013

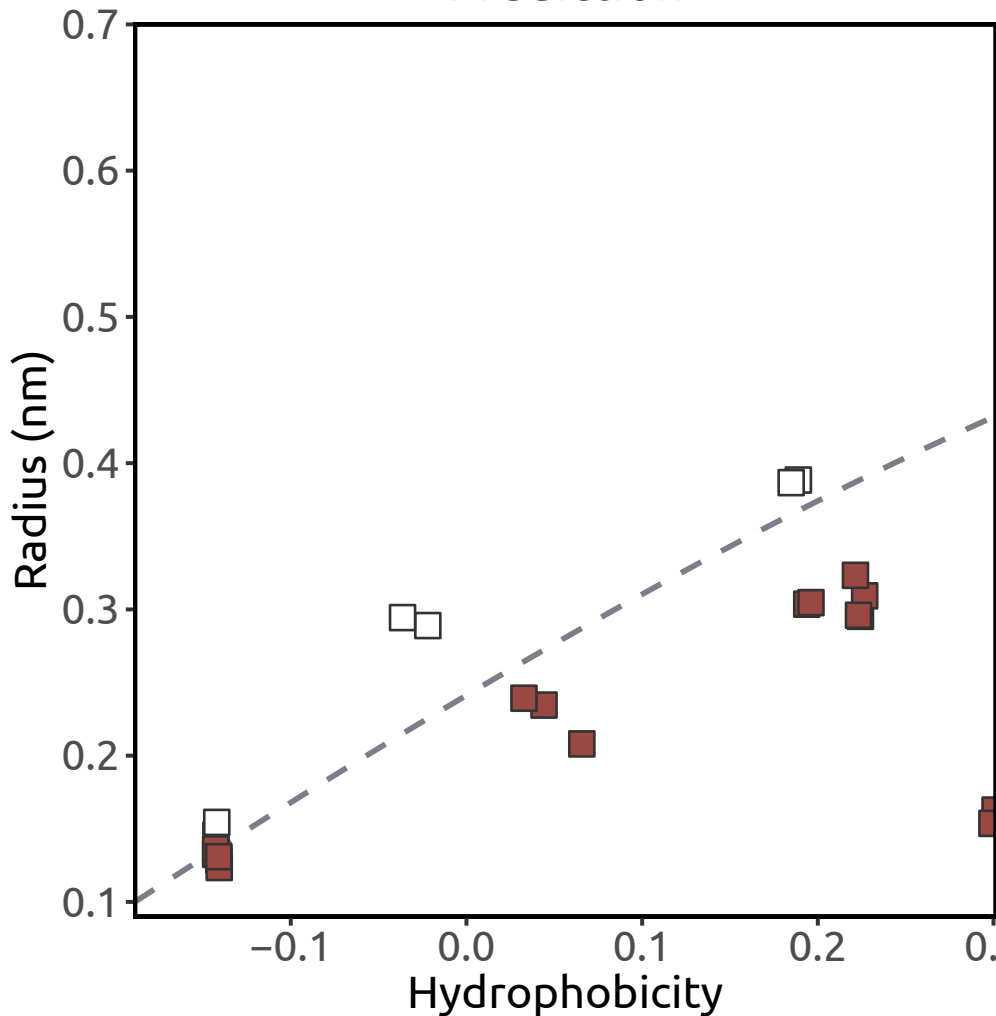
Pore radius



Hydrophobicity



Prediction



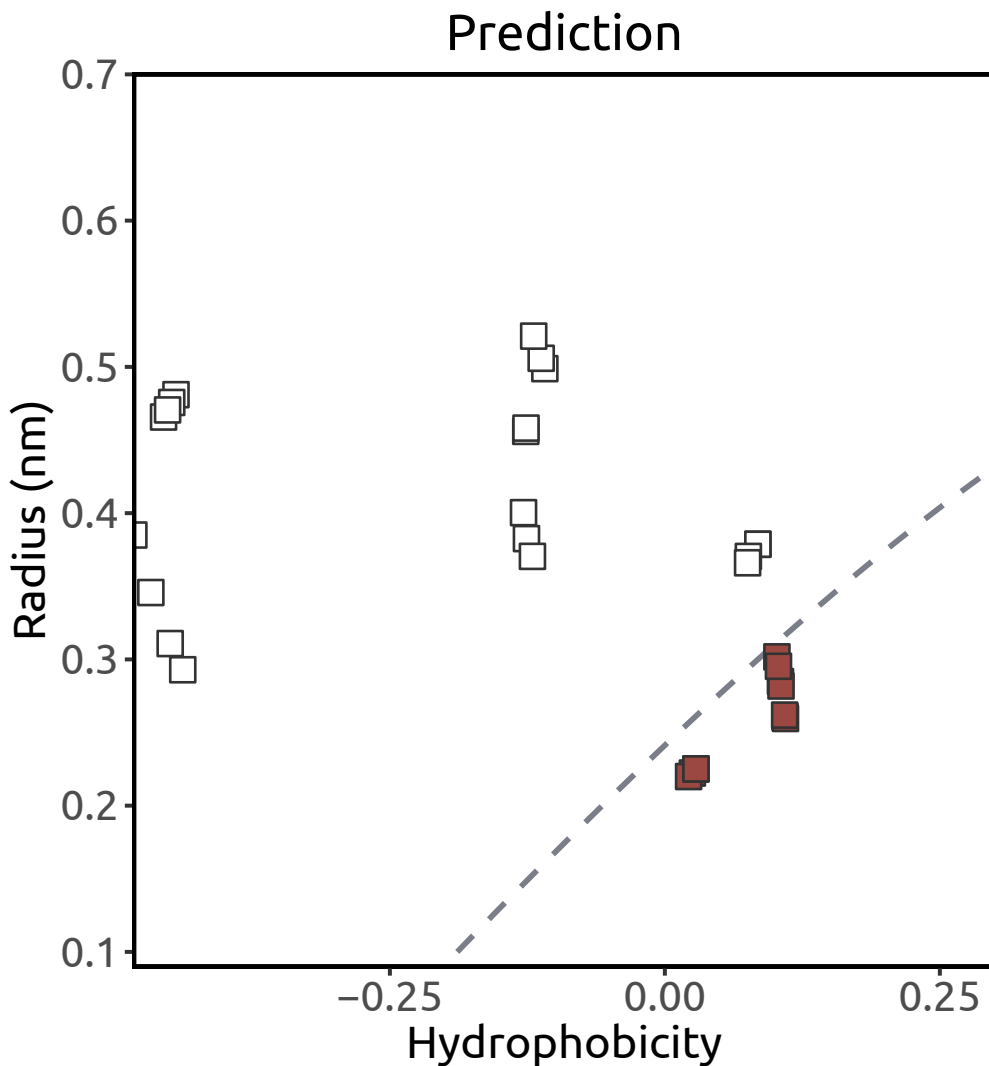
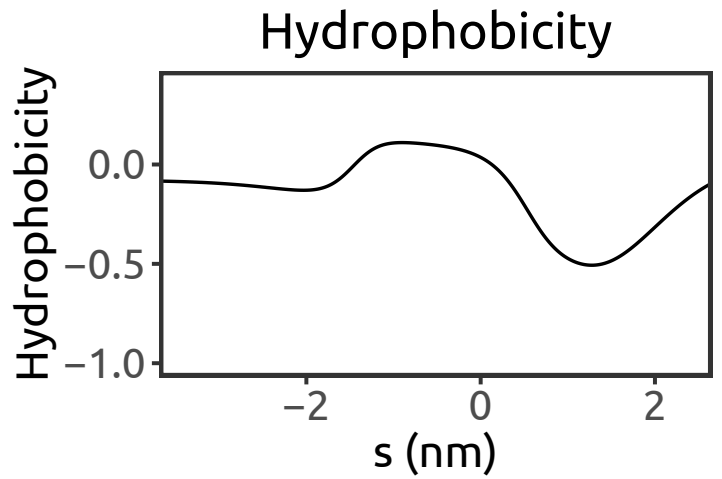
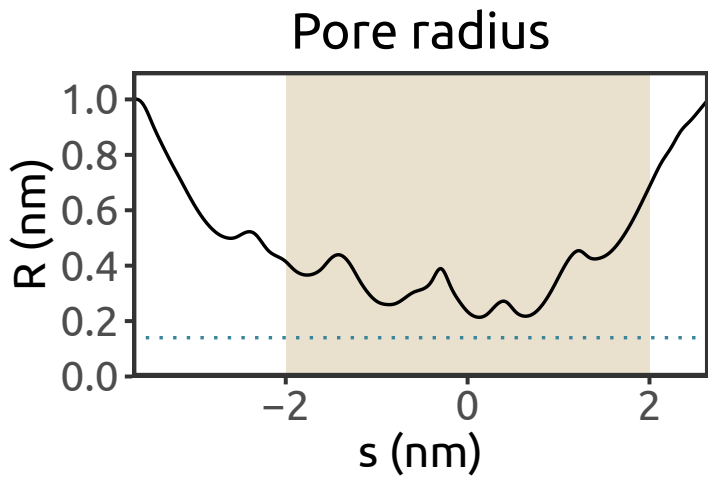
Heuristic score:
1.49 (n = 19)

Simulation result:
barrier to water

TRPV1 (PDB ID: 5IRX)

Rattus norvegicus
cryo-EM (2.95 Å)

Gao et al., 2015



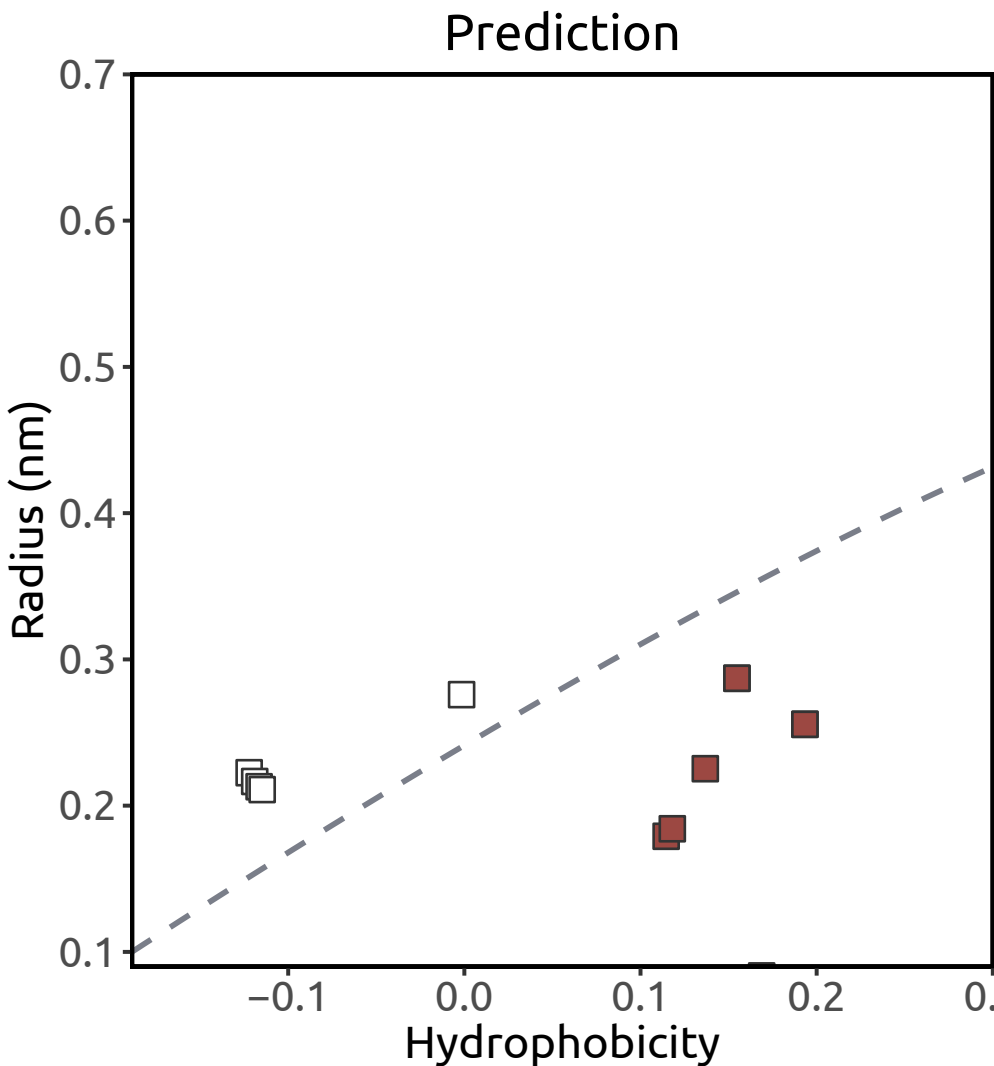
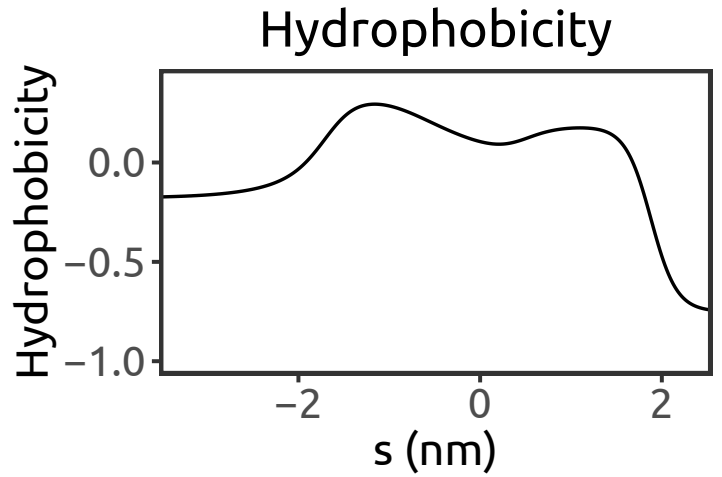
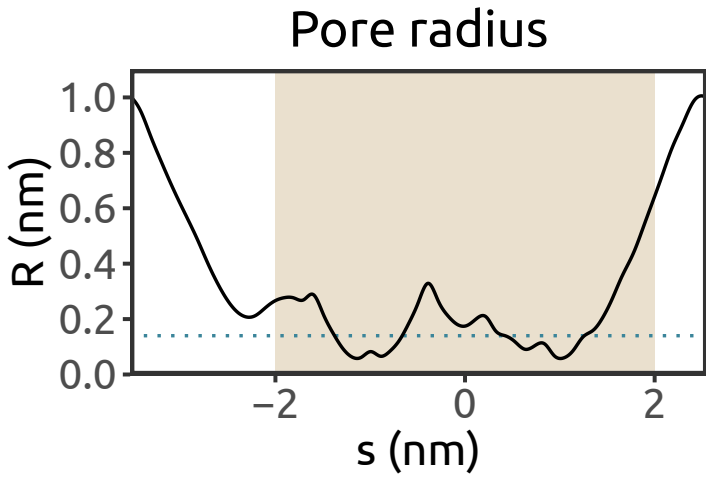
Heuristic score:
0.38 (n = 12)

Simulation result:
hydrated channel

TRPV1 (PDB ID: 5IRZ)

Rattus norvegicus
cryo-EM (3.28 Å)

Gao et al., 2015



Heuristic score:
2.79 (n = 15)

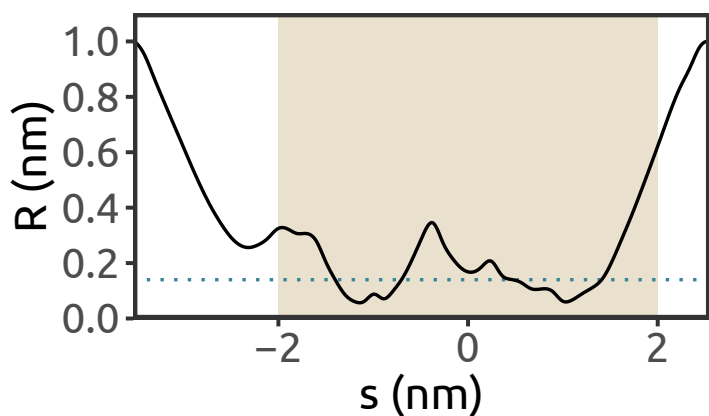
Simulation result:
barrier to water

TRPV1 (PDB ID: 5IS0)

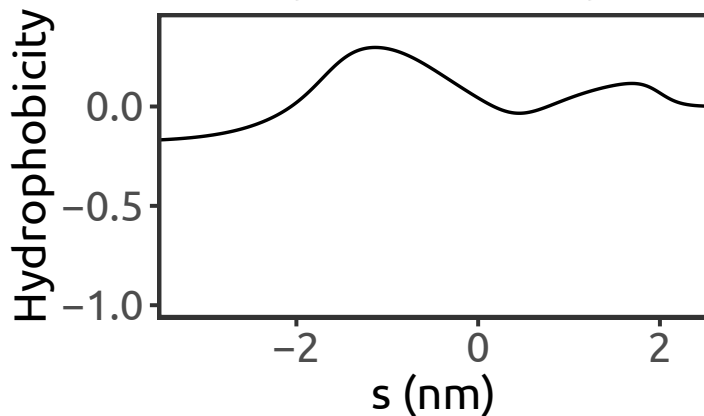
Rattus norvegicus
cryo-EM (3.43 Å)

Gao et al., 2015

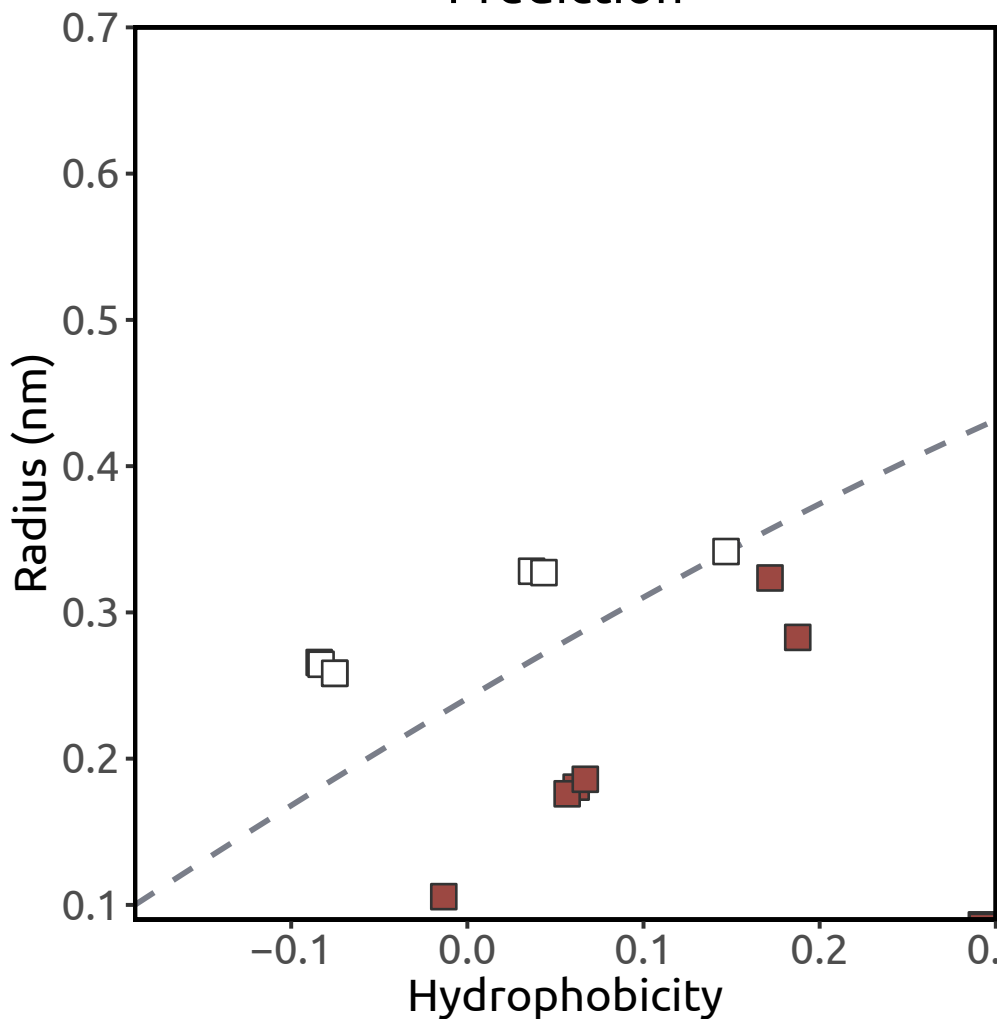
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.16 (n = 13)

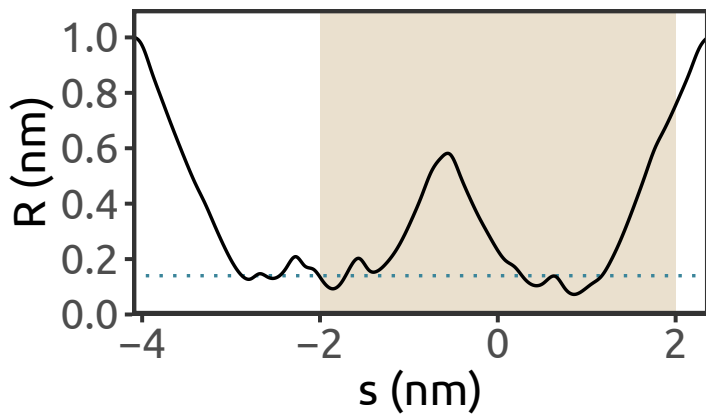
Simulation result:
barrier to water

TRPV2 (PDB ID: 5AN8)

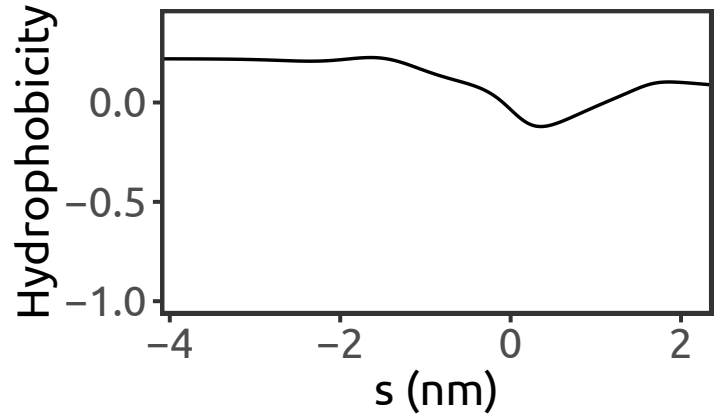
Oryctolagus cuniculus
cryo-EM (3.8 Å)

Zubcevic et al., 2016

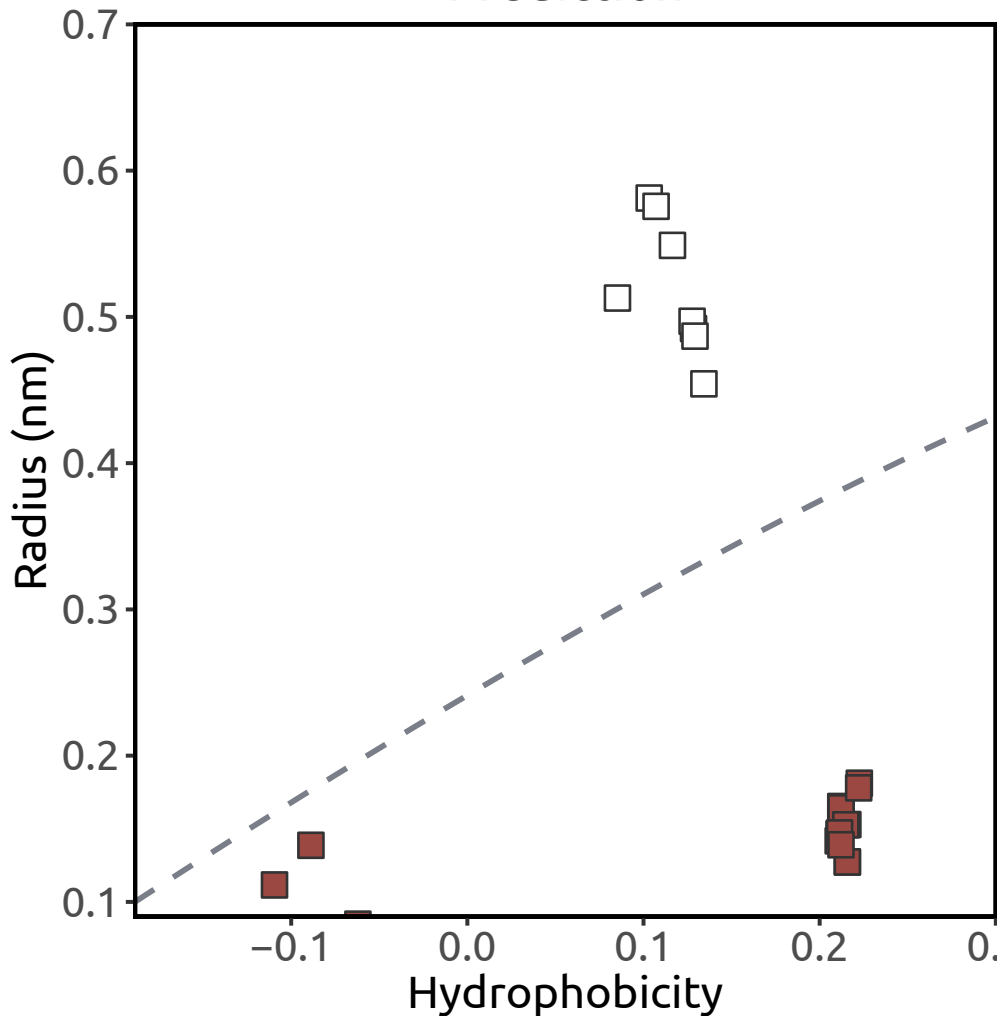
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.72 (n = 17)

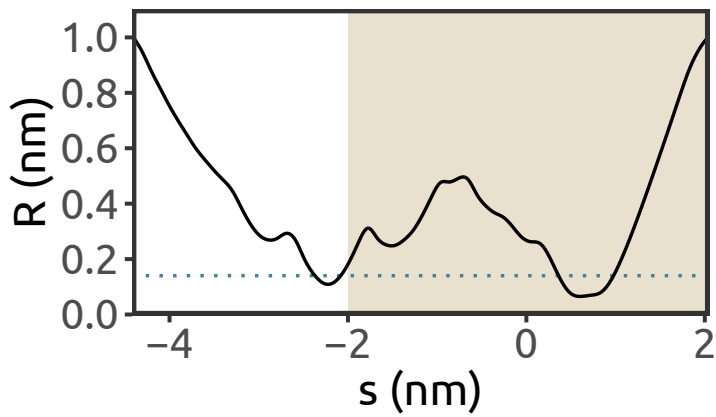
Simulation result:
barrier to water

TRPV2 (PDB ID: 5HI9)

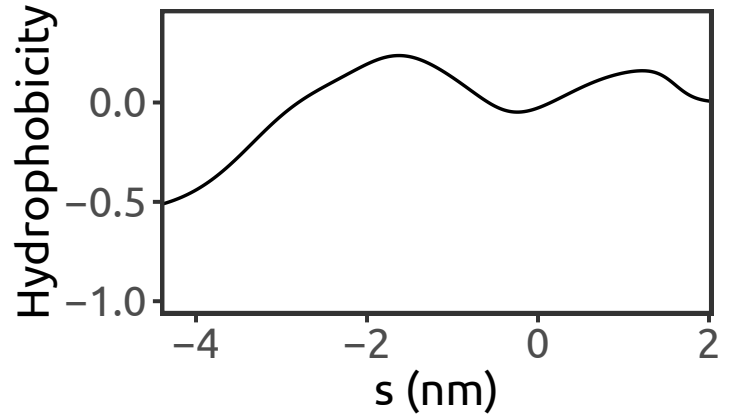
Rattus norvegicus
cryo-EM (4.4 Å)

Huynh et al., 2016

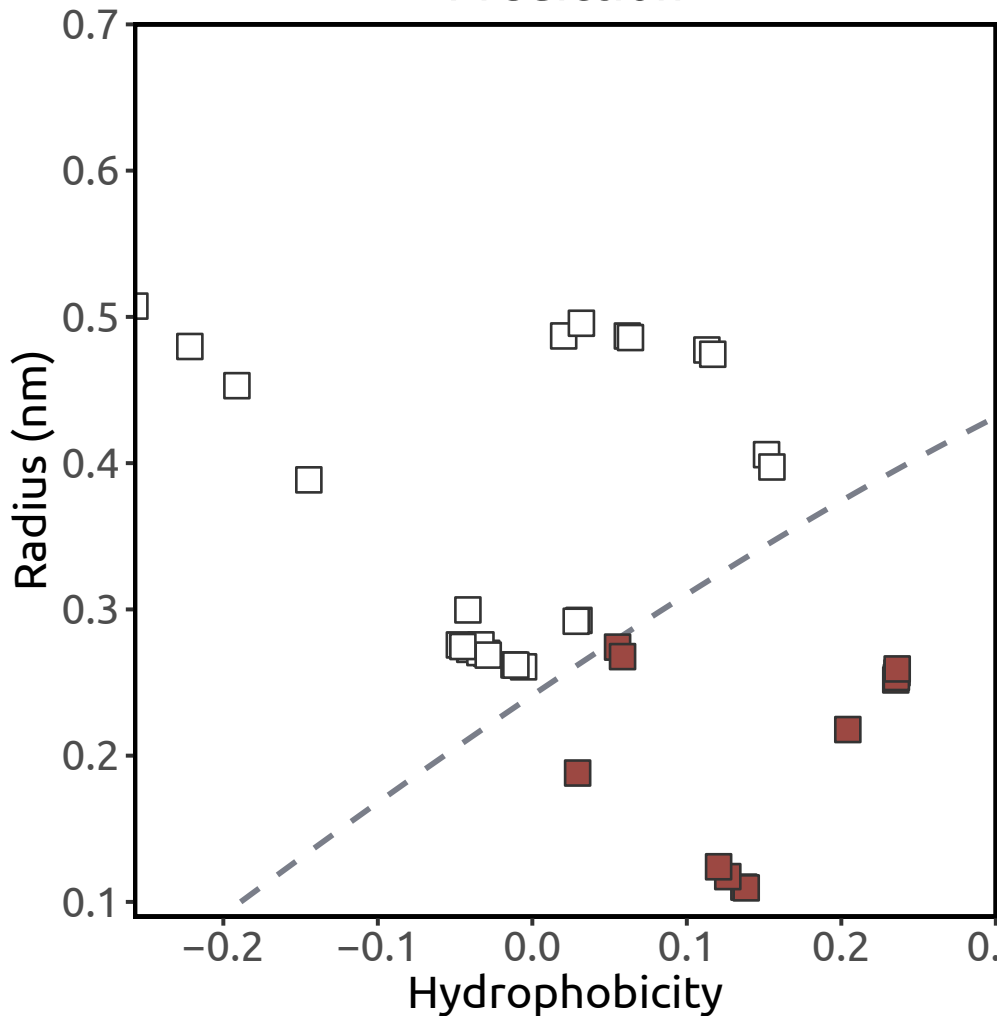
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.17 (n = 16)

Simulation result:
barrier to water

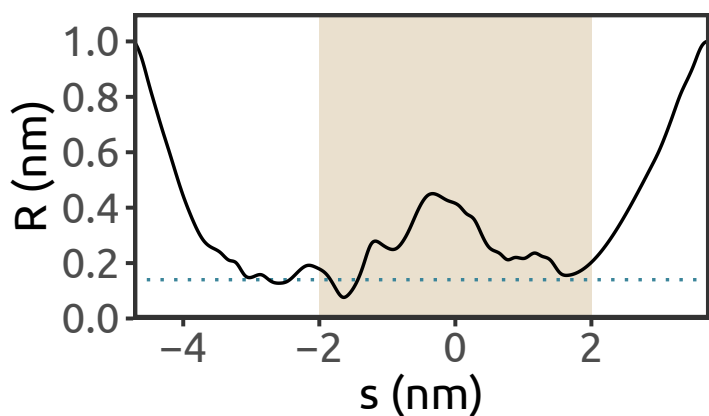
TRPV2 (PDB ID: 6BWJ)

Oryctolagus cuniculus

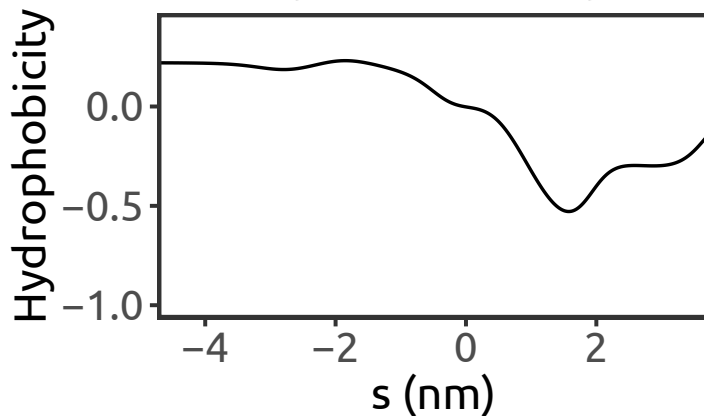
X-ray (3.1 Å)

Zubcevic et al., 2018

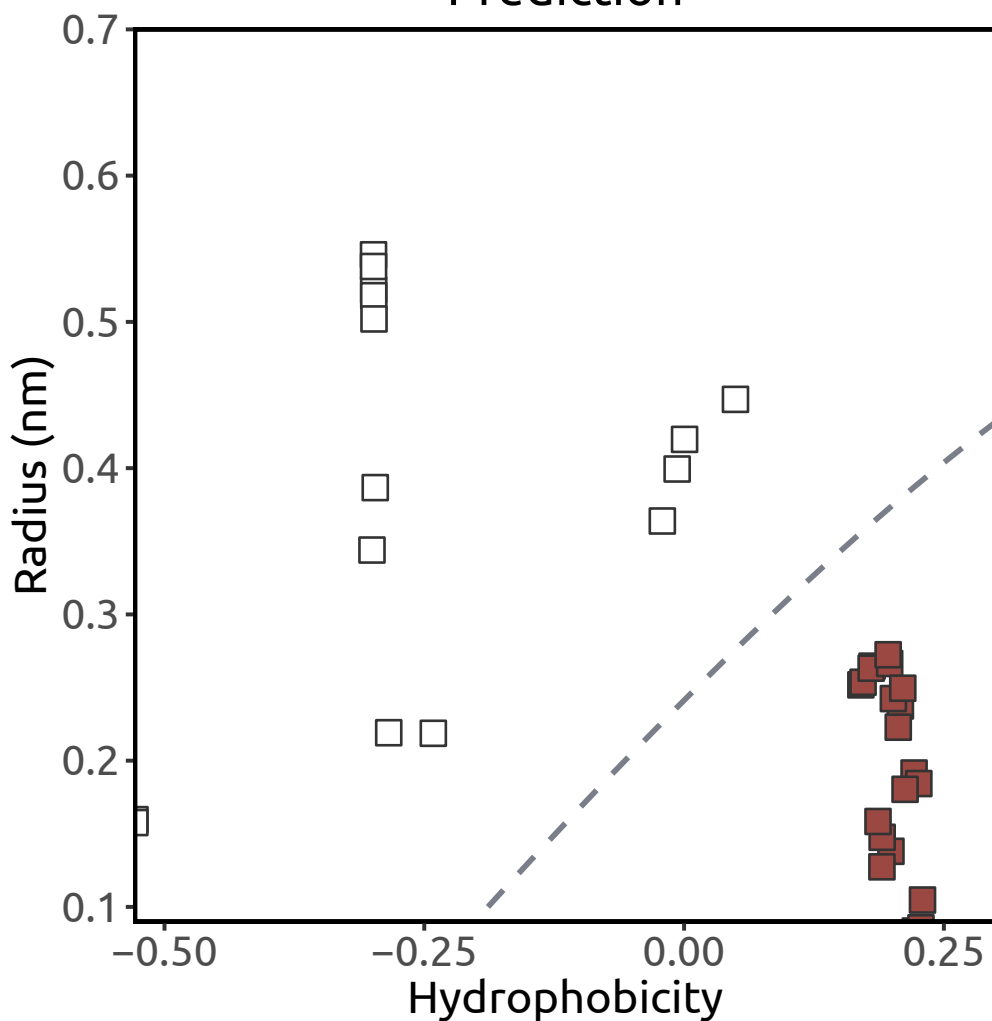
Pore radius



Hydrophobicity



Prediction



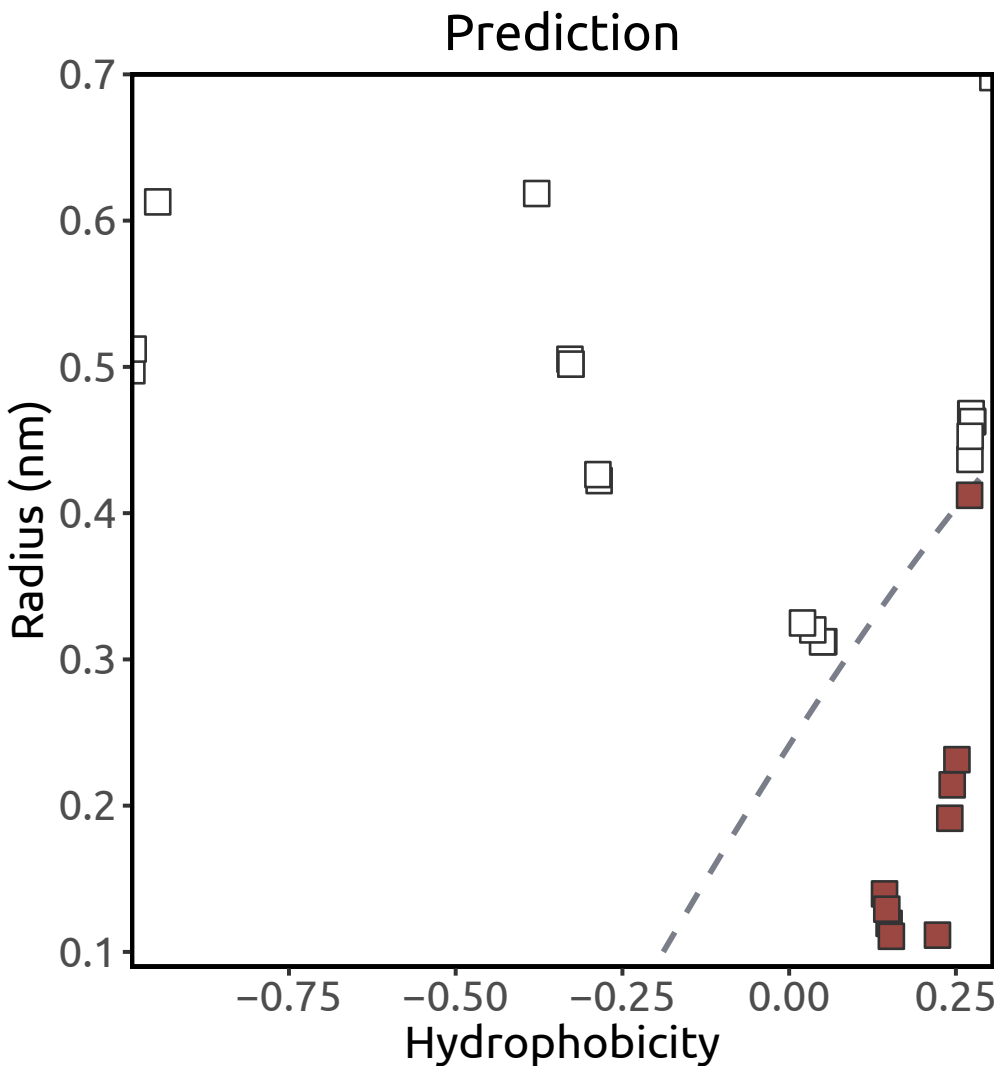
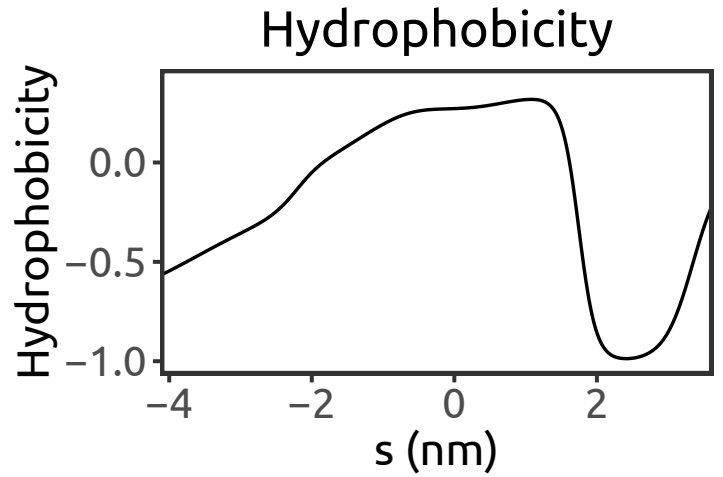
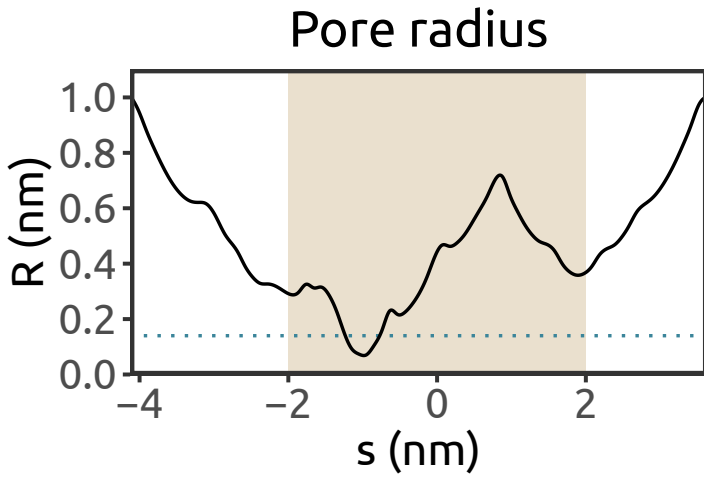
Heuristic score:
3.02 (n = 20)

Simulation result:
barrier to water

TRPV4 (PDB ID: 6BBJ)

Xenopus tropicalis
cryo-EM (3.8 Å)

Deng et al., 2018



Heuristic score:
1.44 (n = 9)

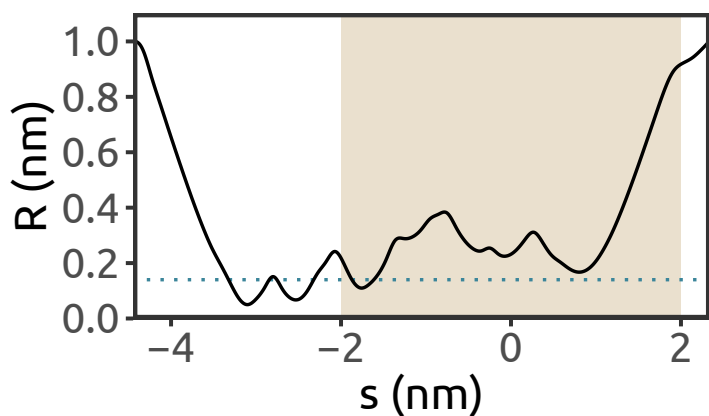
Simulation result:
barrier to water

TRPV5 (PDB ID: 6B5V)

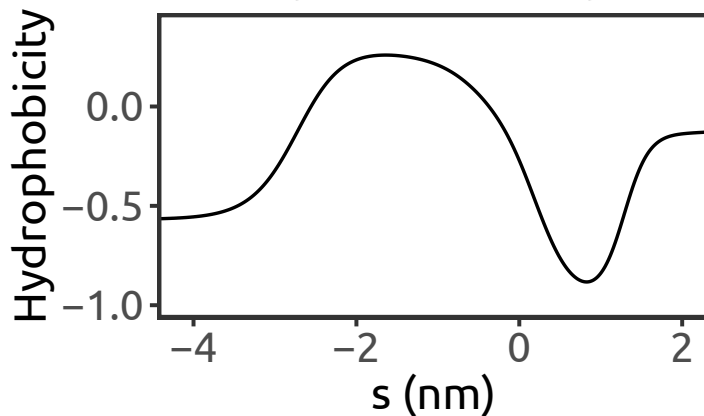
Oryctolagus cuniculus
cryo-EM (4.8 Å)

Hughes et al., 2018

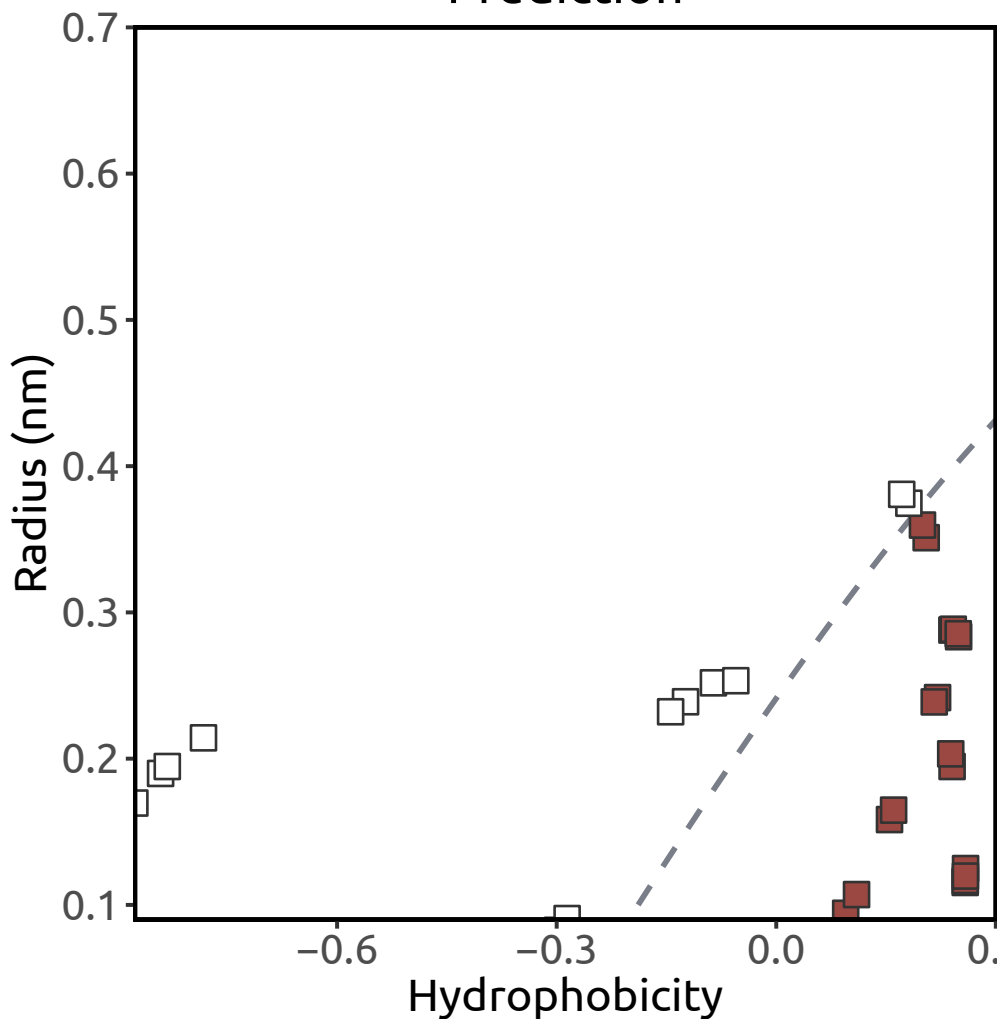
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.66 (n = 18)

Simulation result:
barrier to water

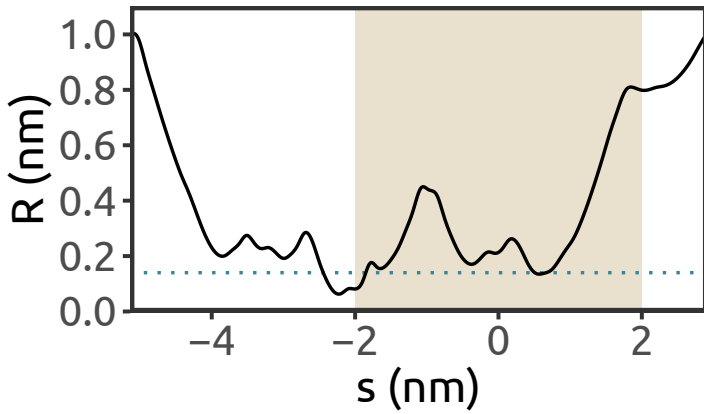
TRPV6 (PDB ID: 5IWK)

Rattus norvegicus

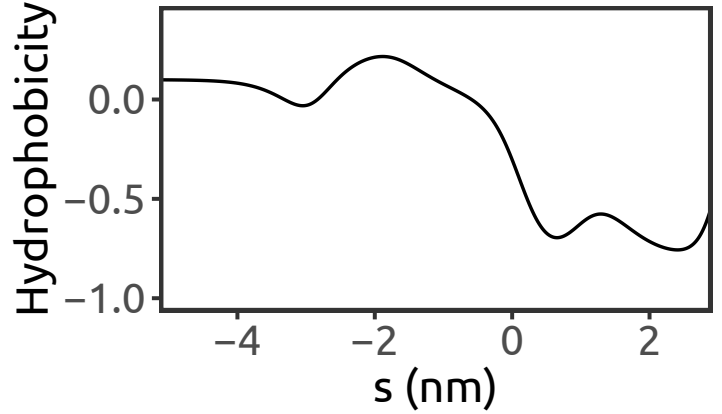
X-ray (3.25 Å)

Saotome et al., 2016

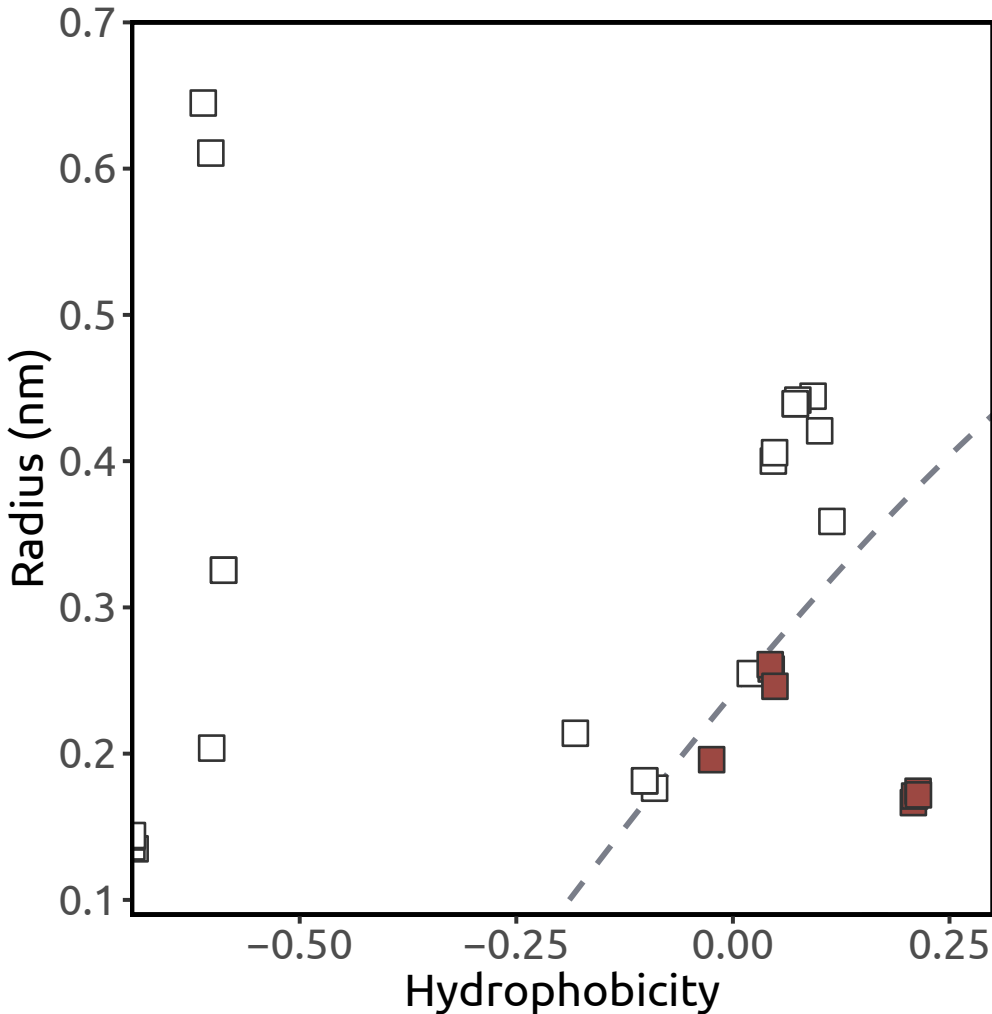
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.71 ($n = 12$)

Simulation result:

barrier to water

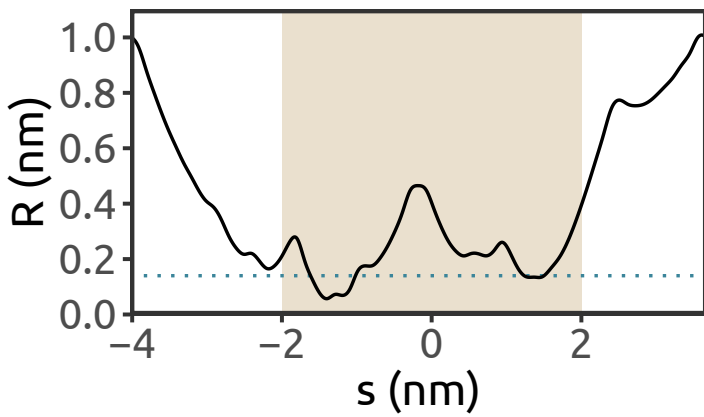
TRPV6 (PDB ID: 5IWP)

Rattus norvegicus

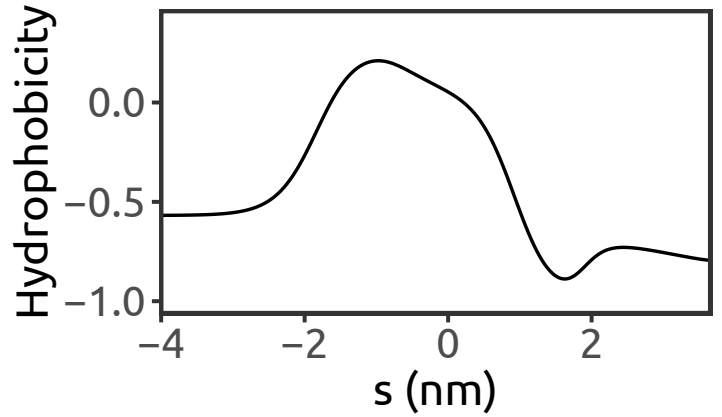
X-ray (3.65 Å)

Saotome et al., 2016

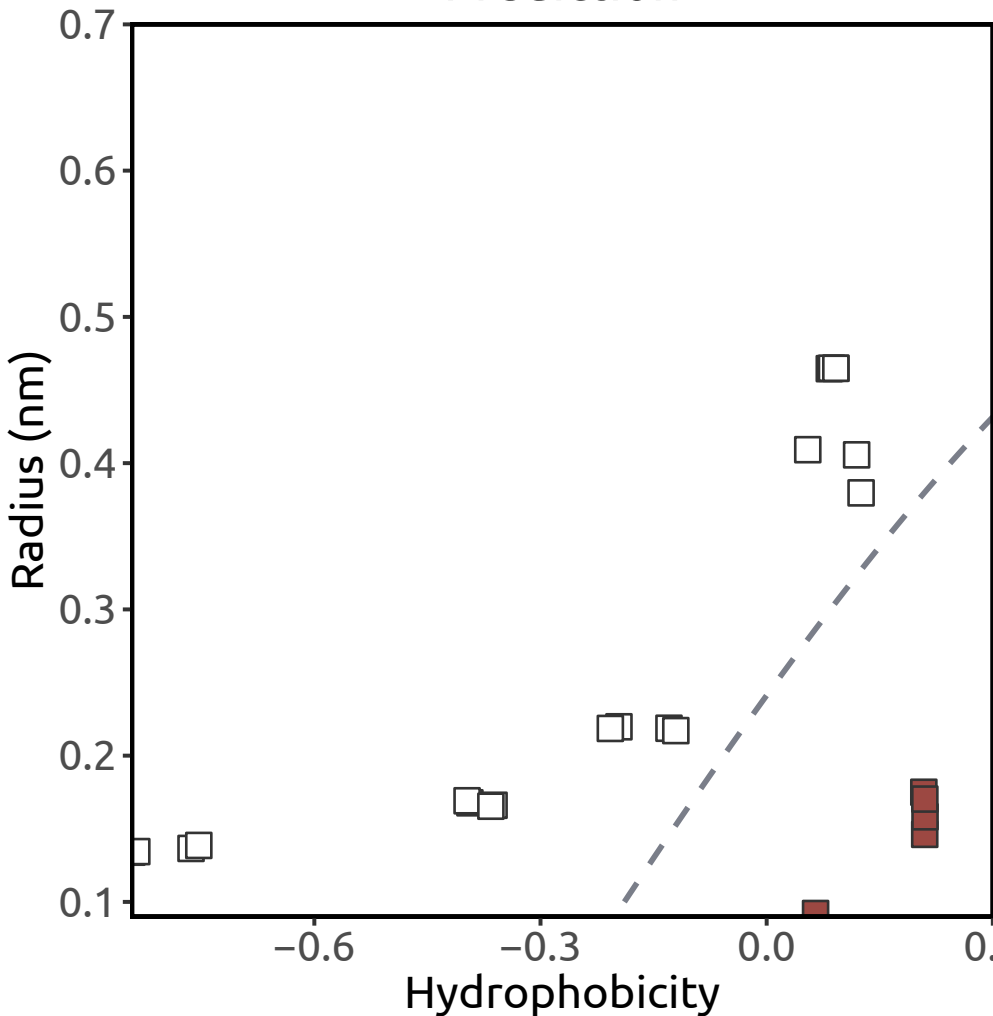
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.47 (n = 8)

Simulation result:

barrier to water

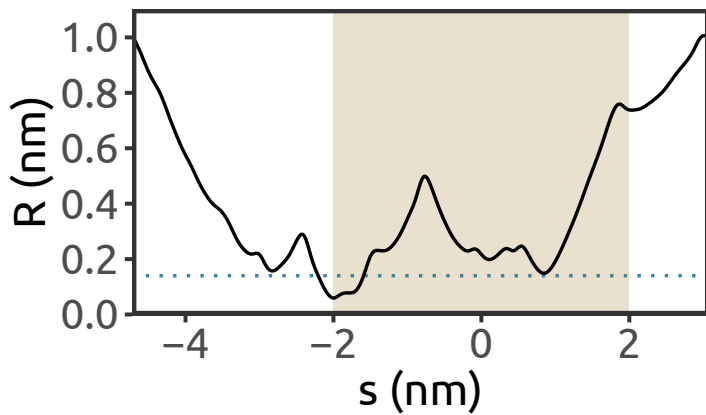
TRPV6 (PDB ID: 5WO6)

Rattus norvegicus

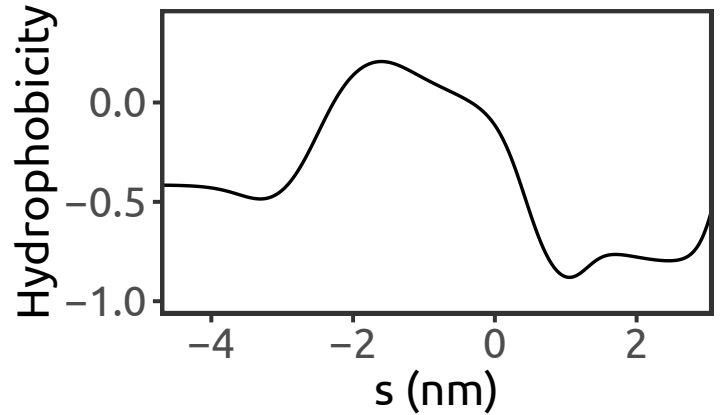
X-ray (3.31 Å)

Singh et al., 2017

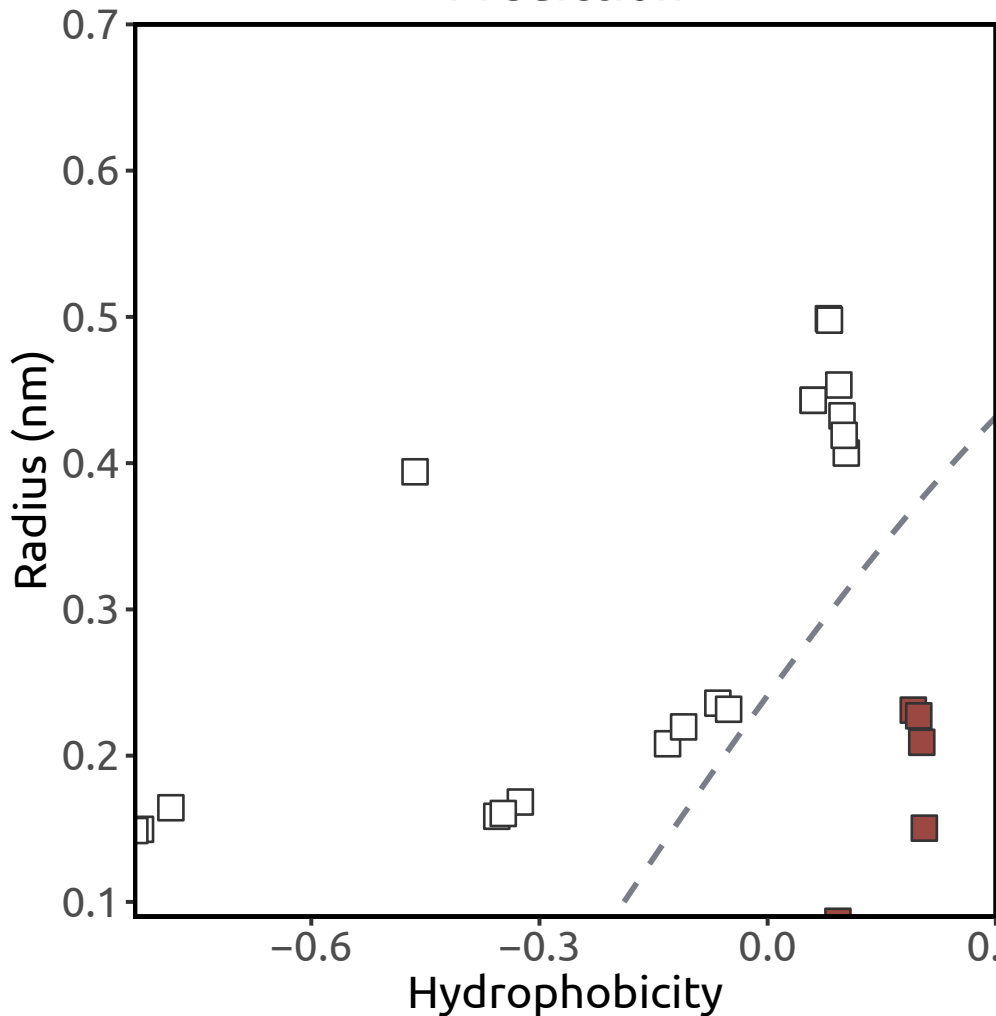
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.34 (n = 8)

Simulation result:
barrier to water

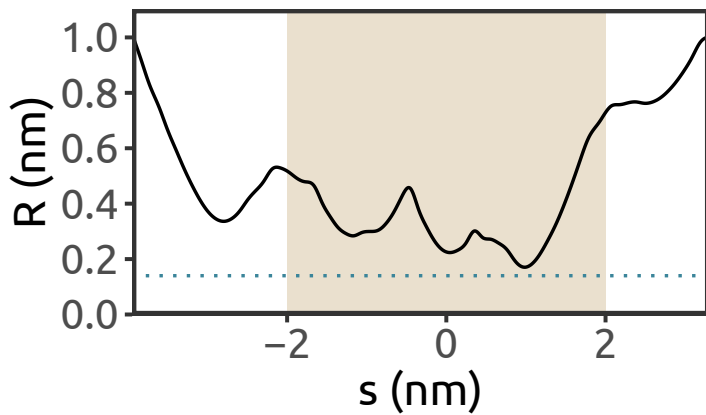
TRPV6 (PDB ID: 6BO8)

Homo sapiens

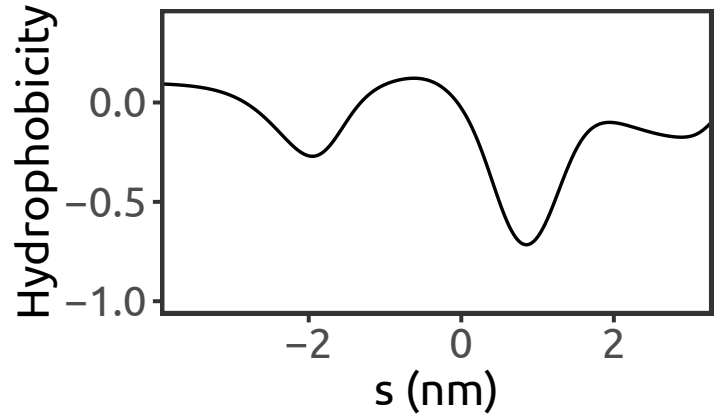
cryo-EM (3.6 Å)

McGoldrick et al., 2018

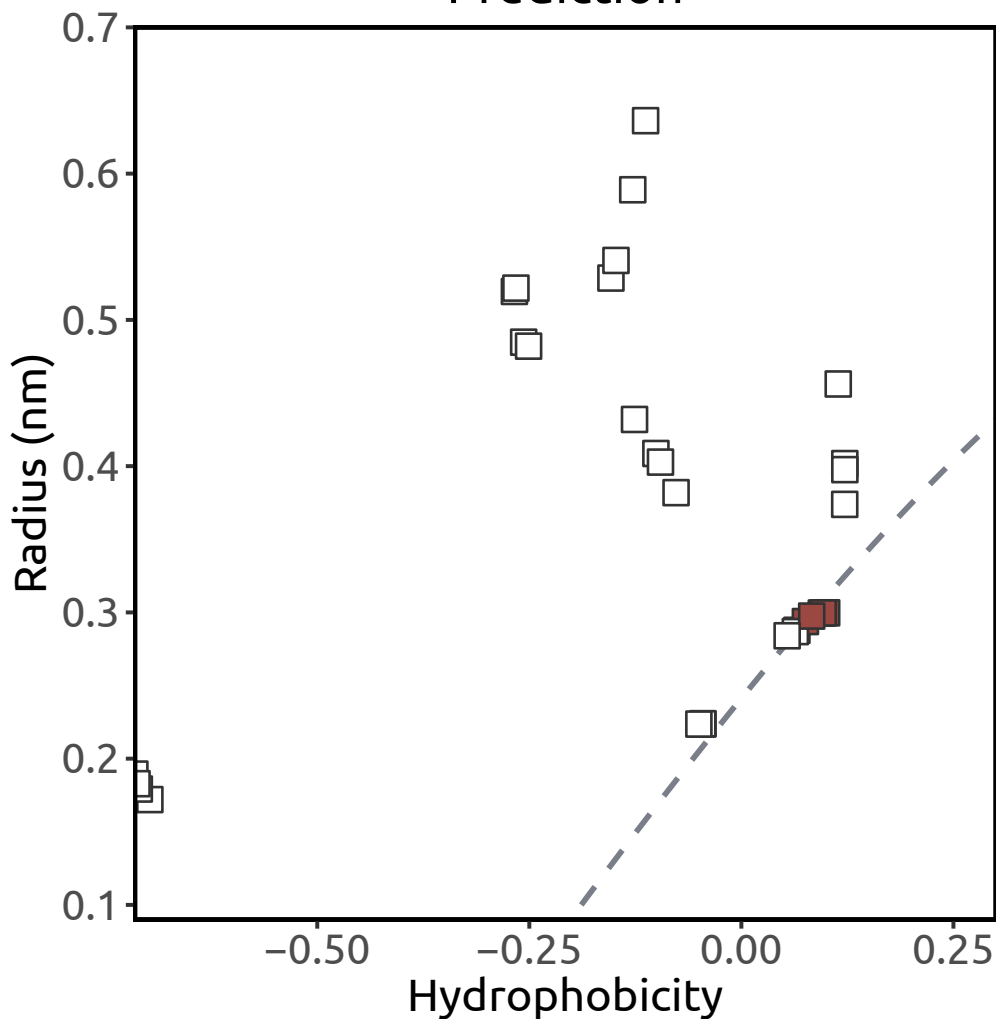
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.03 ($n = 5$)

Simulation result:
hydrated channel

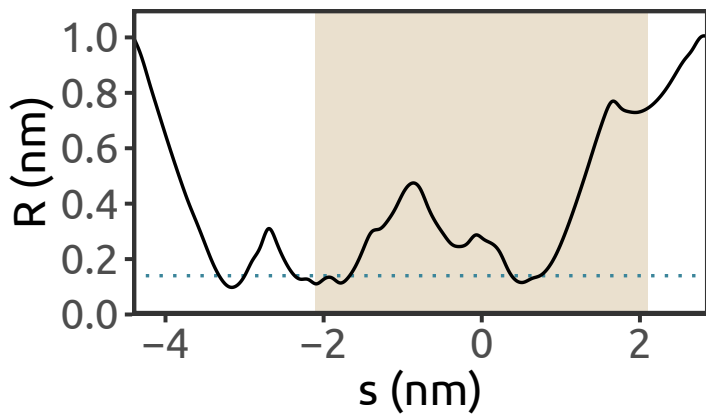
TRPV6 (PDB ID: 6BOB)

Homo sapiens

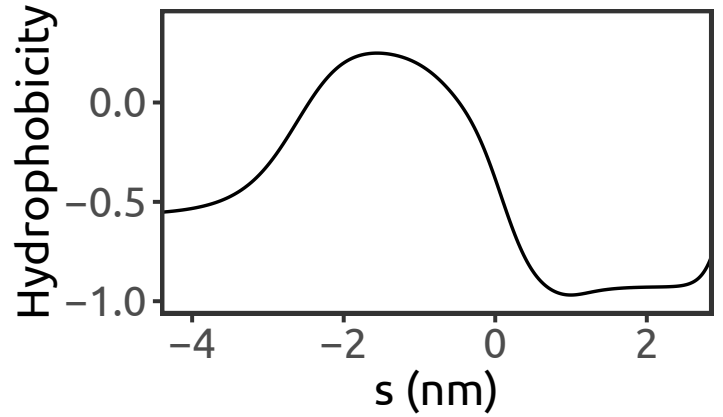
cryo-EM (3.9 Å)

McGoldrick et al., 2018

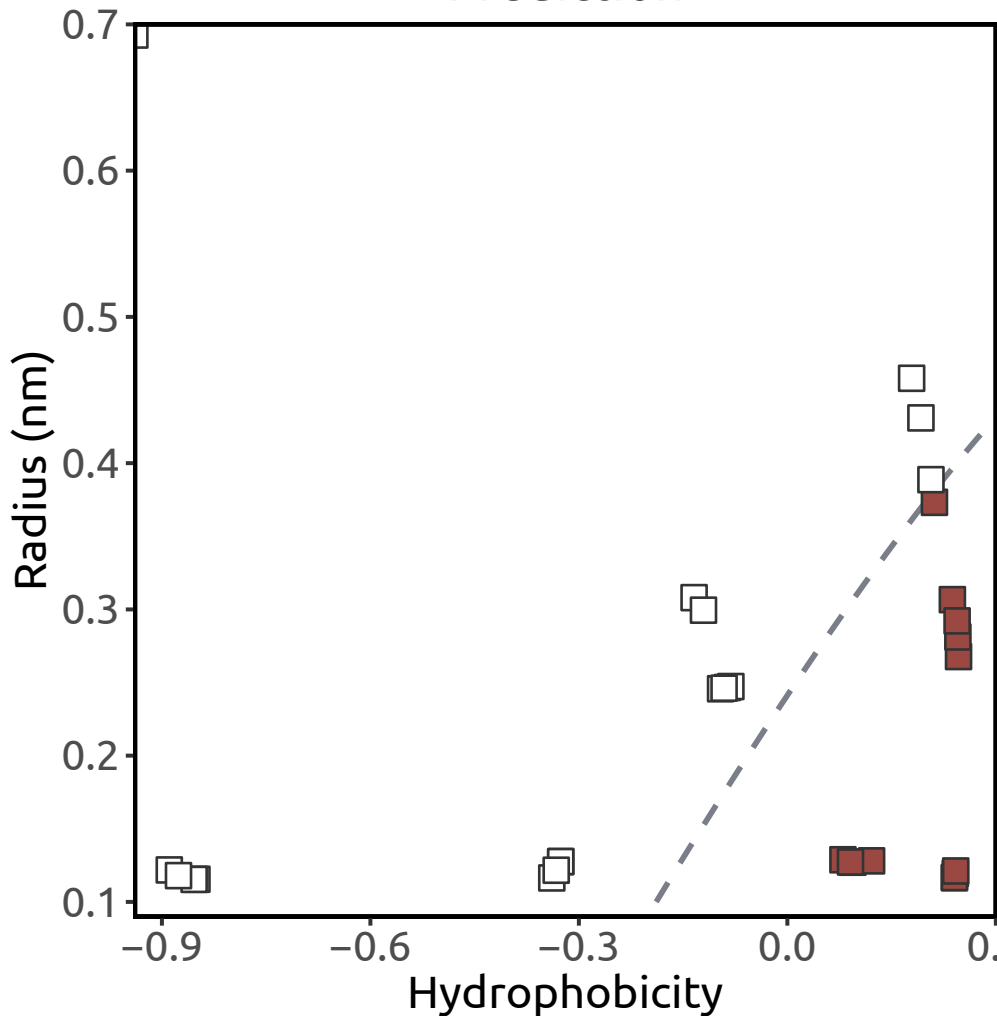
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.94 (n = 13)

Simulation result:

barrier to water

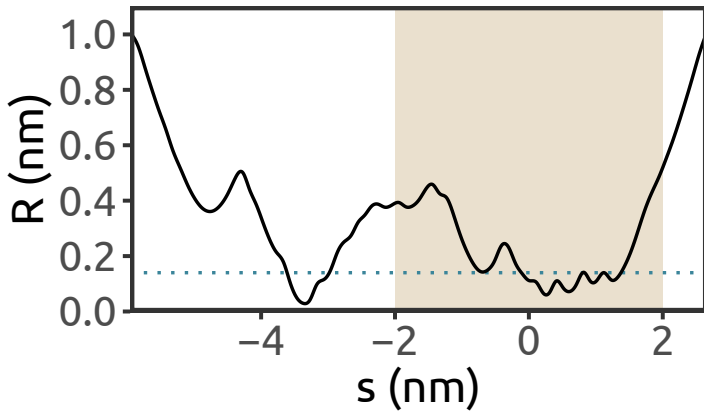
GsuK (PDB ID: 4GX5)

Geobacter sulfurreducens

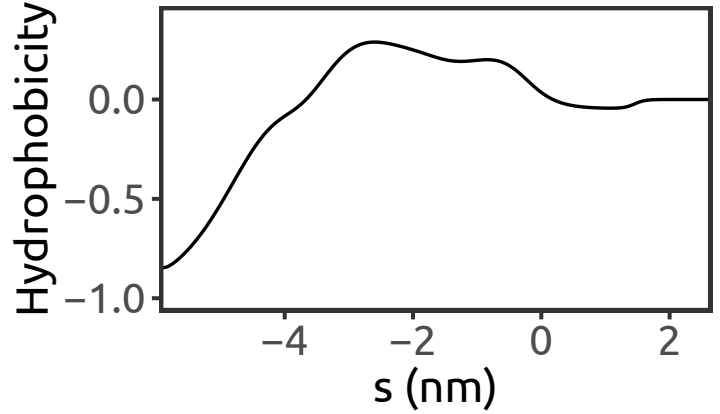
X-ray (3.7 Å)

Kong et al., 2012

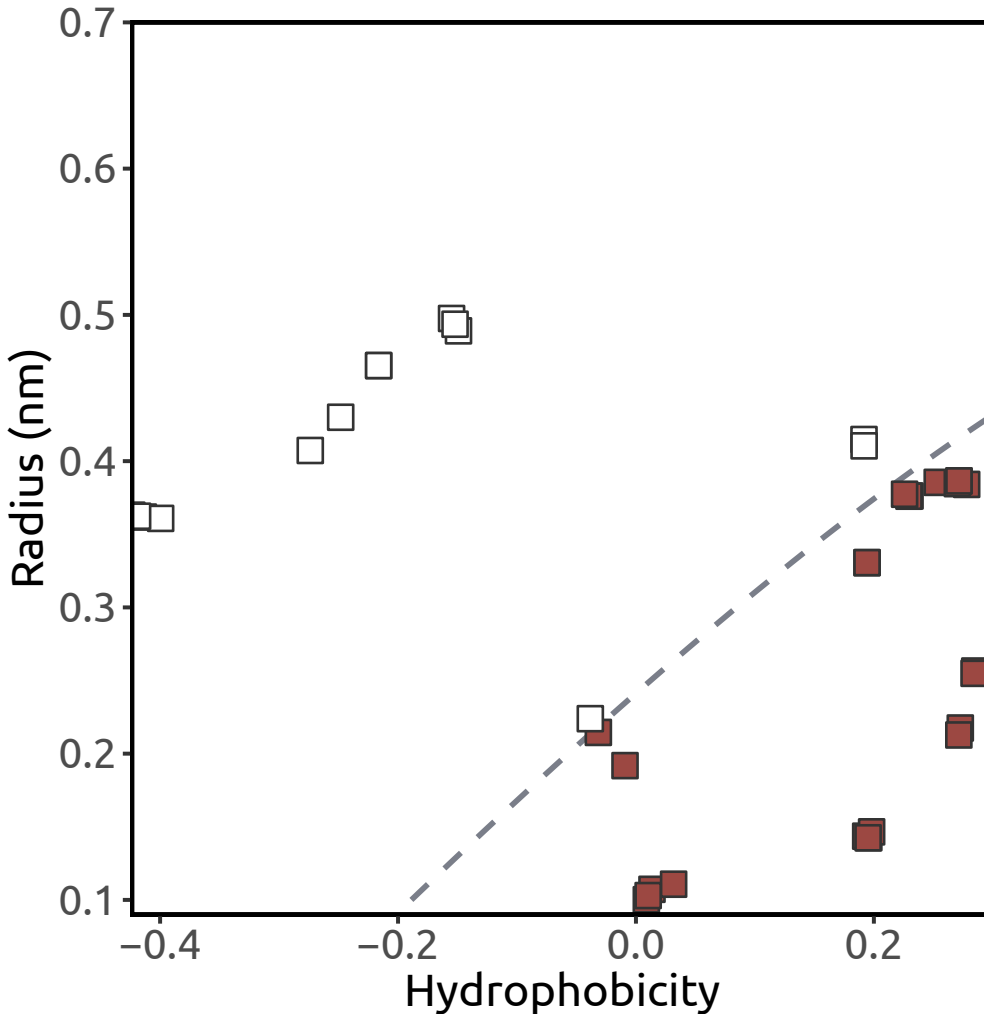
Pore radius



Hydrophobicity



Prediction



Heuristic score:
3.15 (n = 28)

Simulation result:
barrier to water

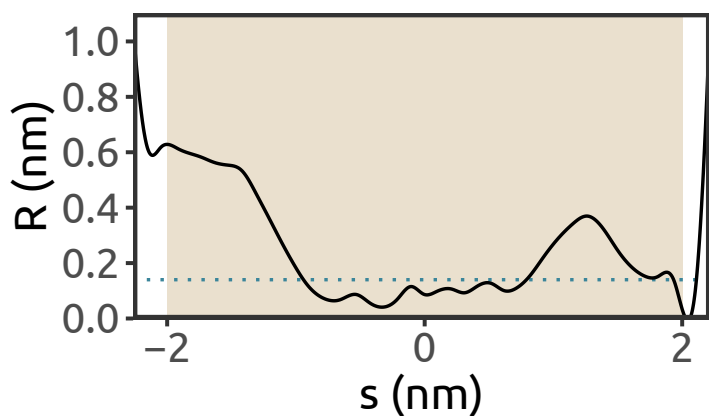
K2P10.1 (PDB ID: 4BW5)

Homo sapiens

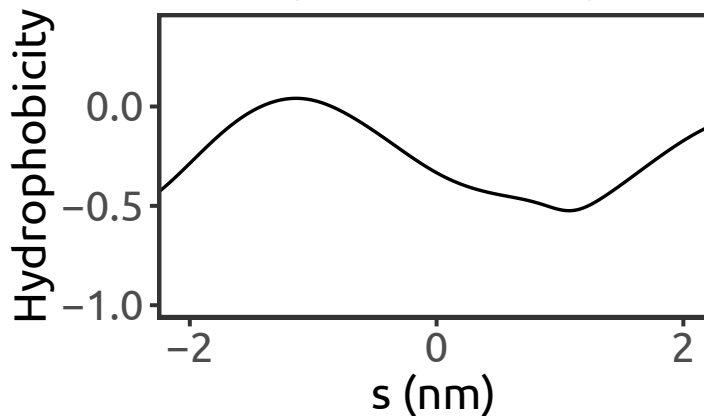
X-ray (3.2 Å)

Dong et al., 2015

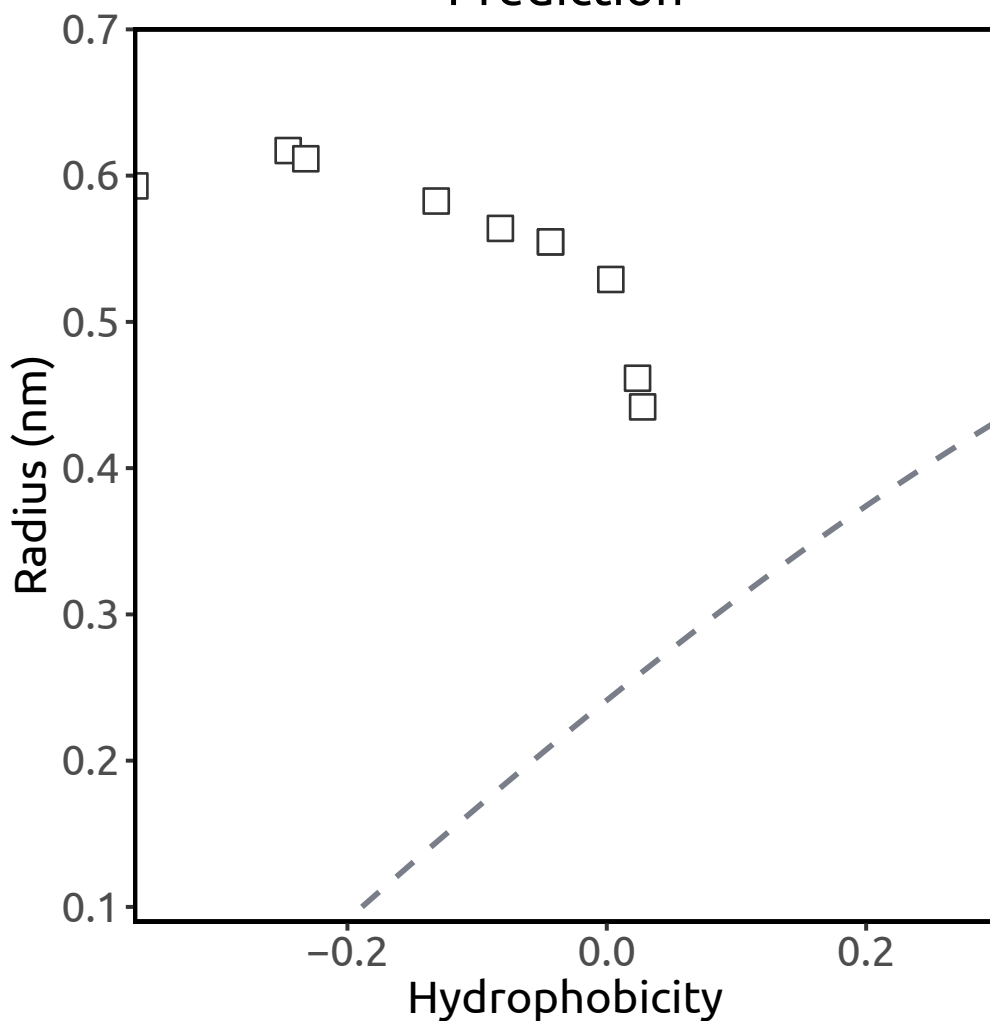
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0 (n = 0)

Simulation result:

hydrated channel

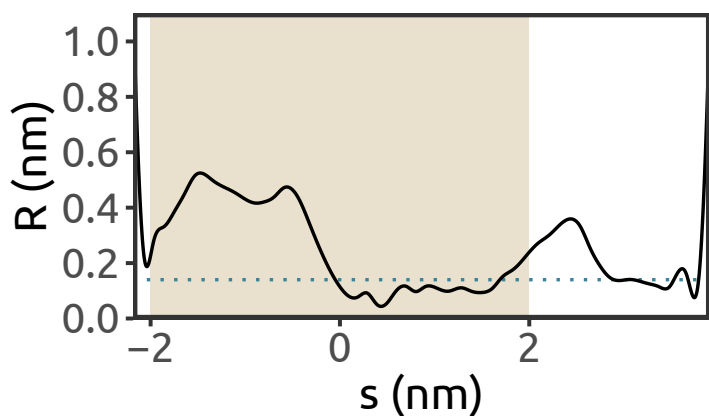
K2P10.1 (PDB ID: 4XDL)

Homo sapiens

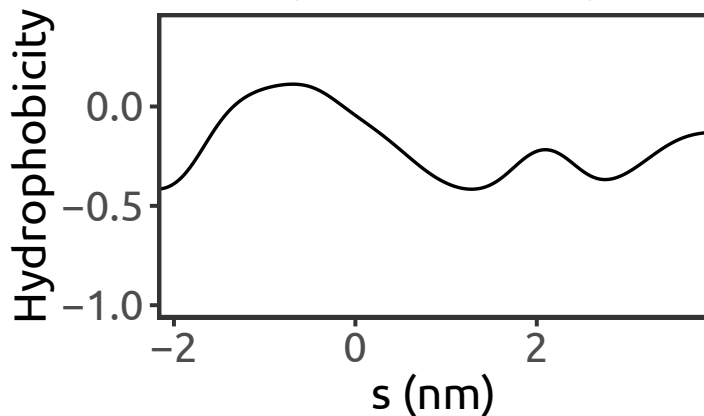
X-ray (3.5 Å)

Dong et al., 2015

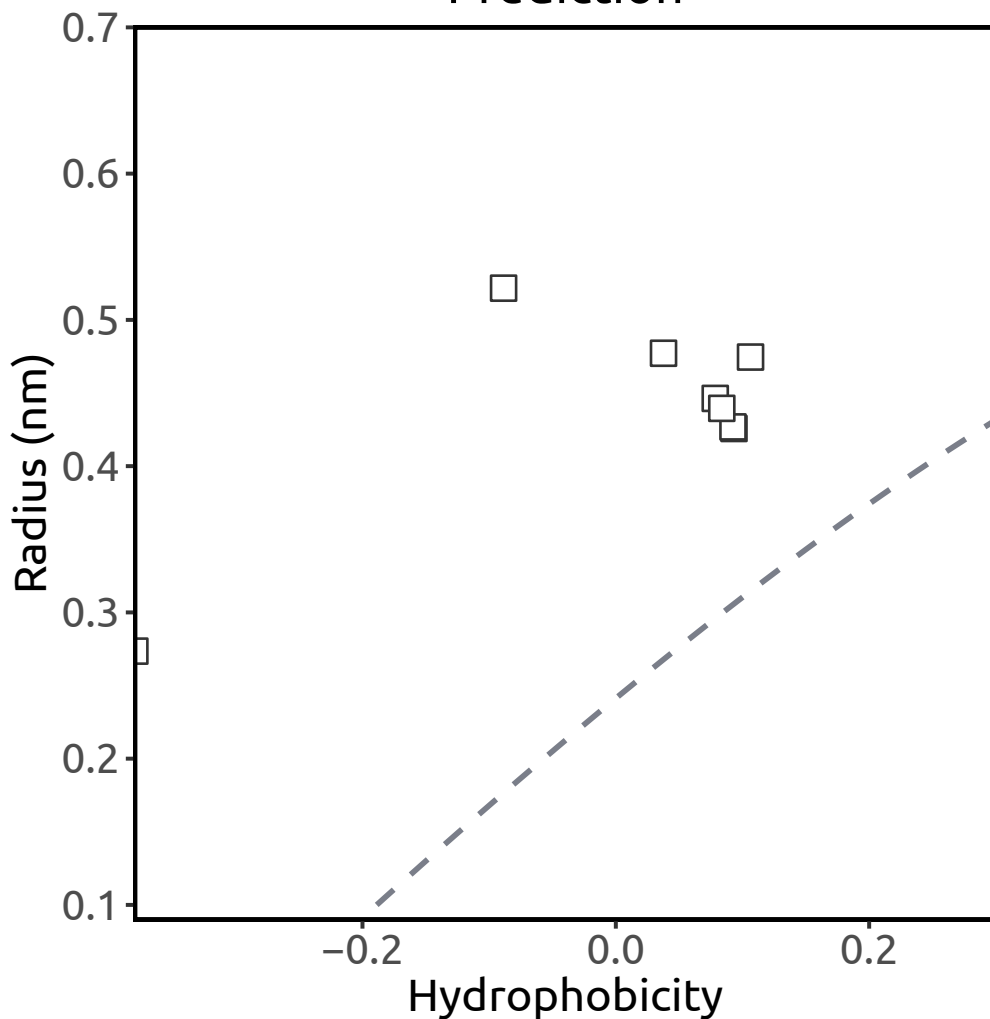
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

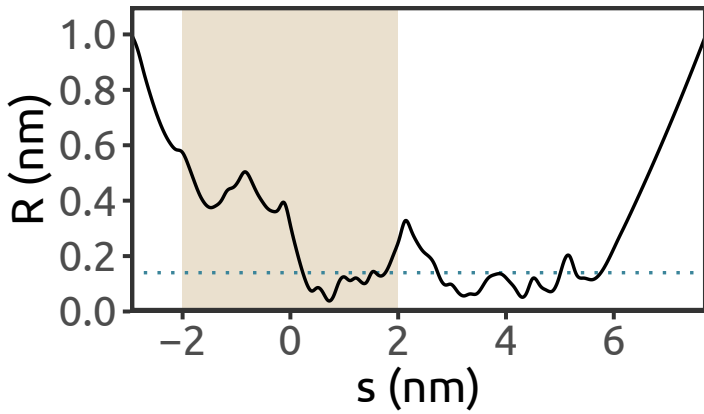
K2P1.1 (PDB ID: 3UKM)

Homo sapiens

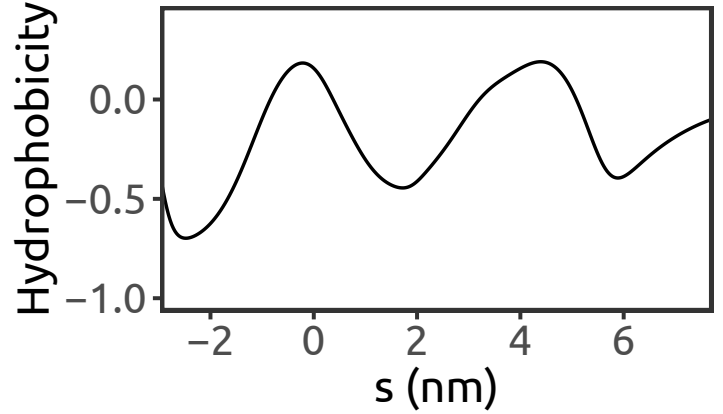
X-ray (3.4 Å)

Miller & Long, 2012

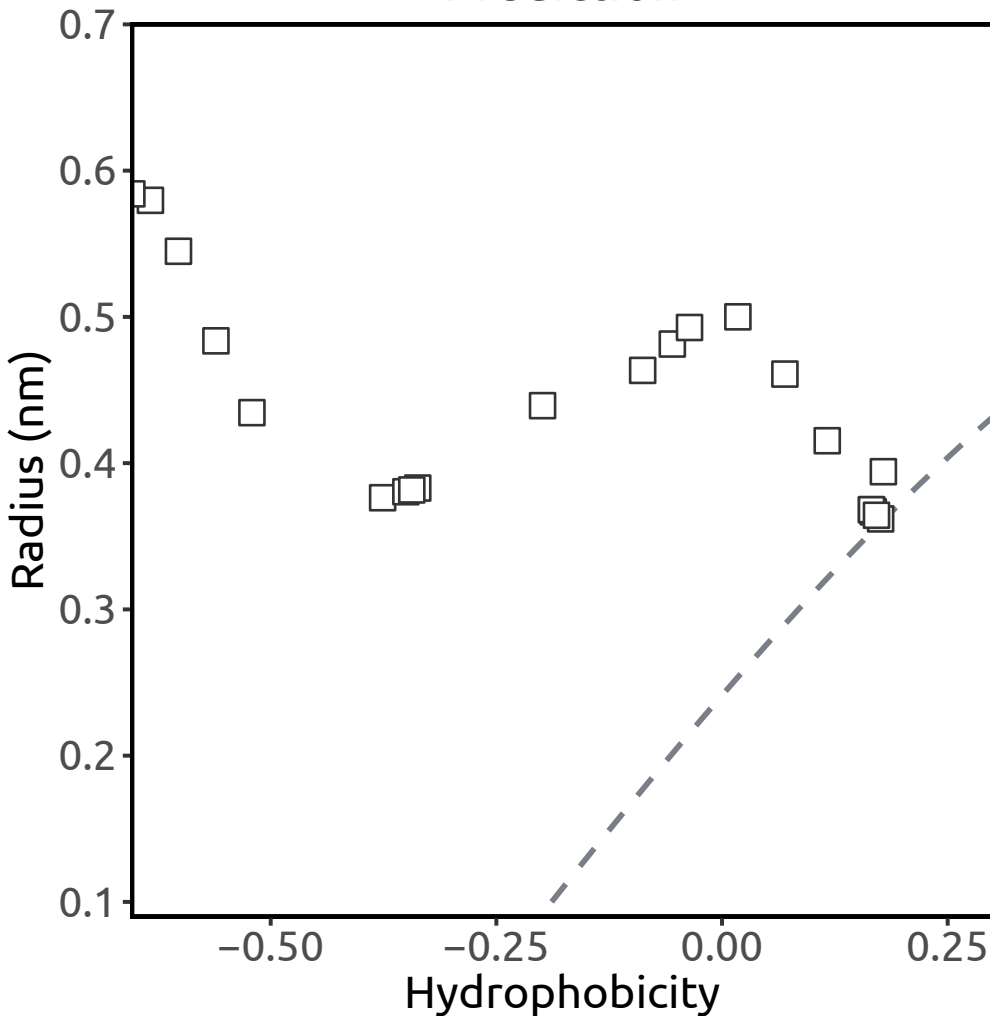
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0 (n = 0)

Simulation result:

hydrated channel

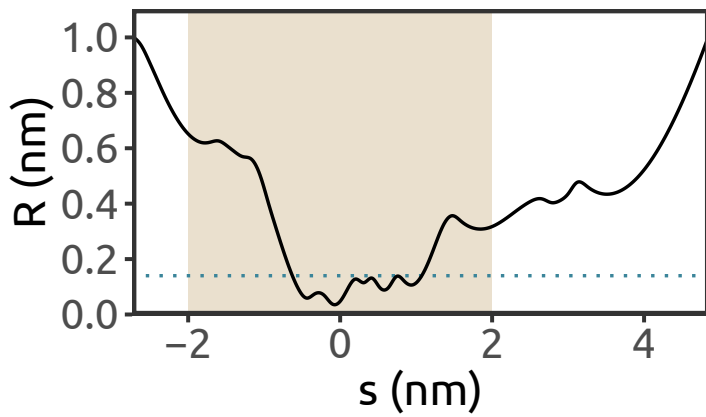
K2P2.1 (PDB ID: 5VKP)

Mus musculus

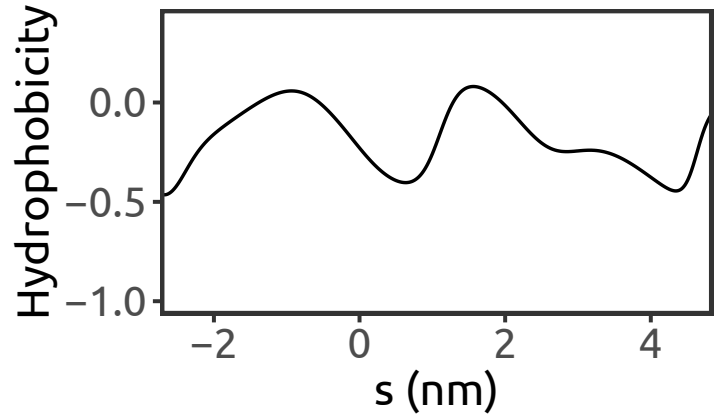
X-ray (2.8 Å)

Lolicato et al., 2017

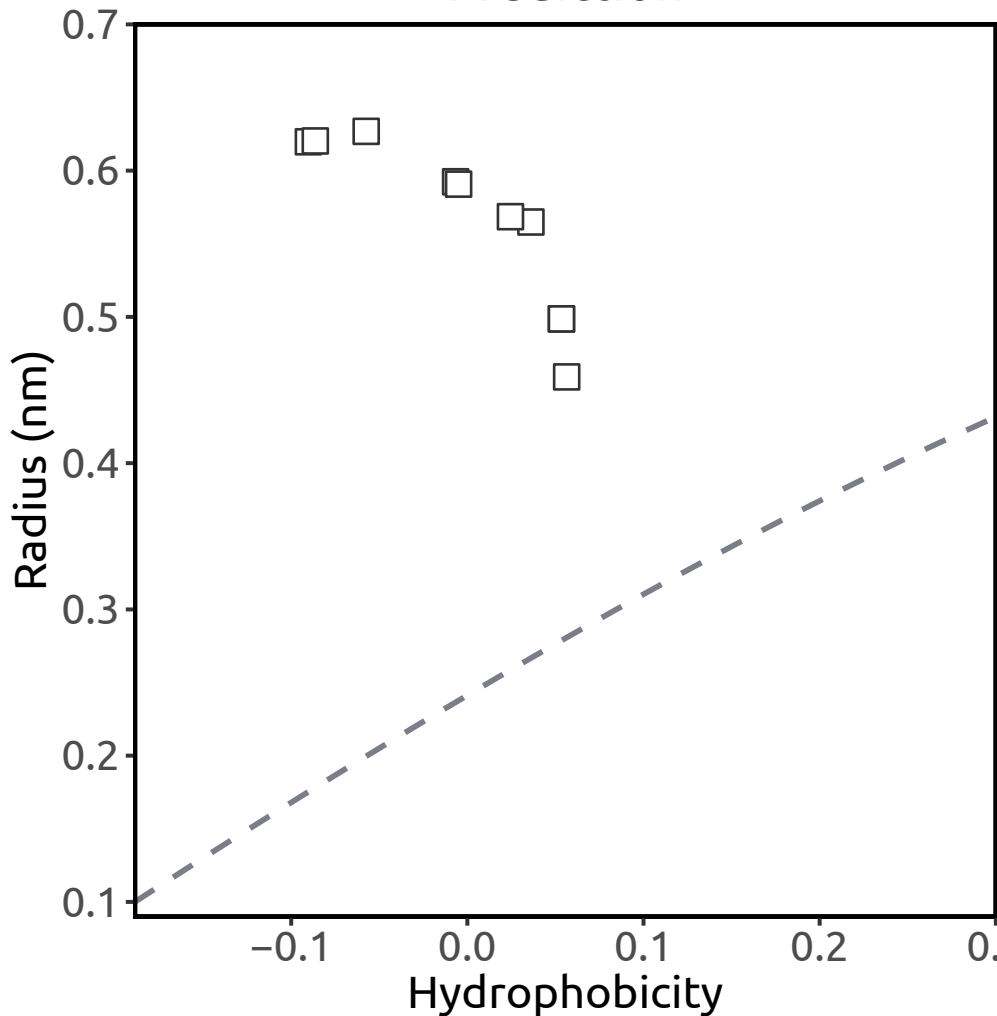
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

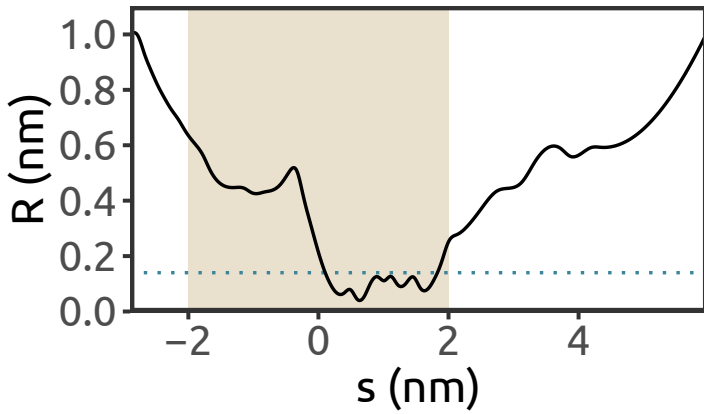
K2P4.1 (PDB ID: 4WFE)

Homo sapiens

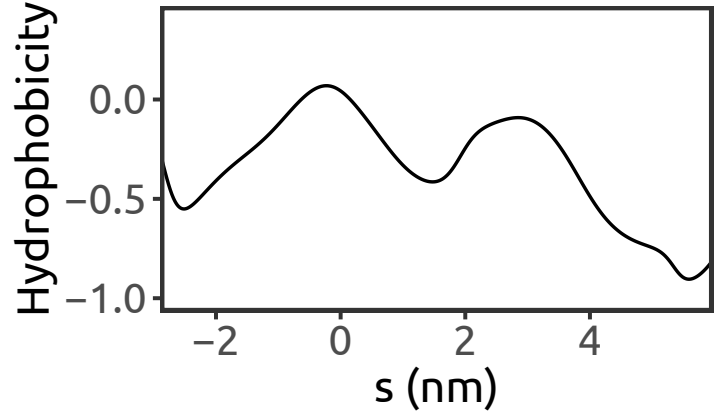
X-ray (2.5 Å)

Brohawn et al., 2014

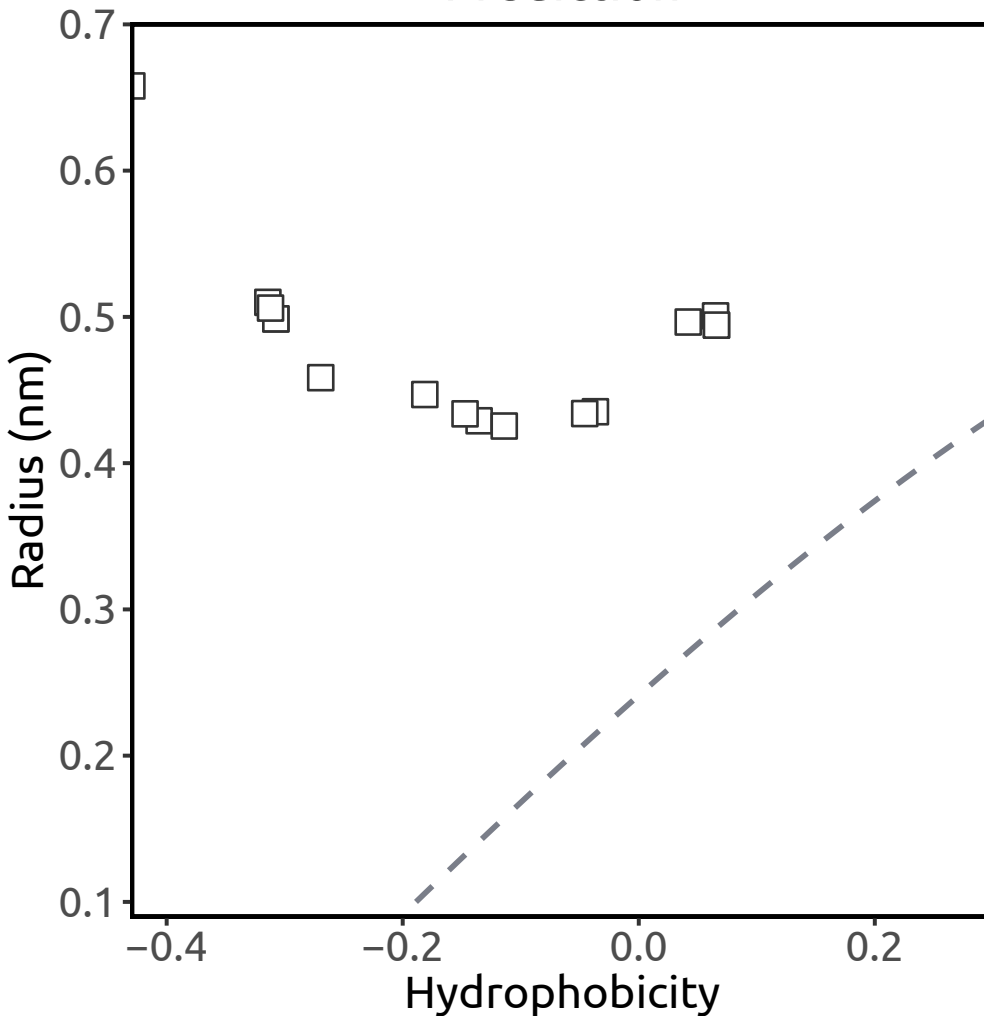
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 ($n = 0$)

Simulation result:
hydrated channel

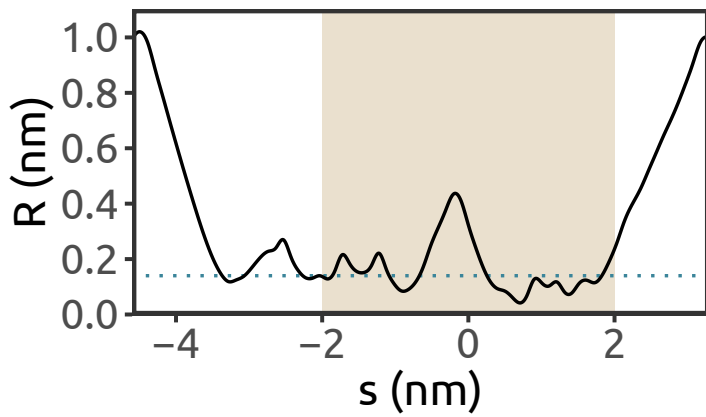
KcsA (PDB ID: 1K4C)

Streptomyces lividans

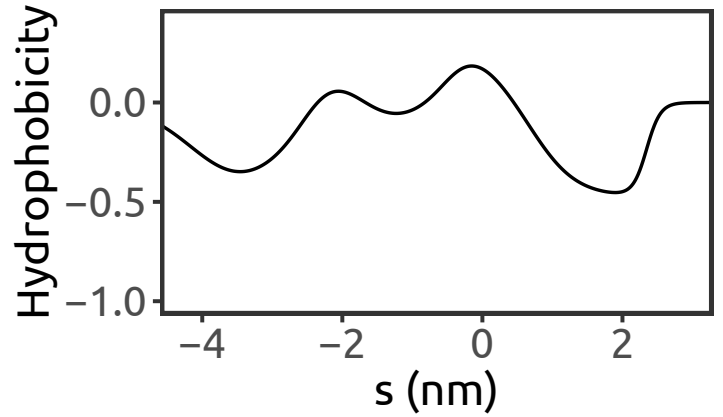
X-ray (2 Å)

Zhou et al., 2001

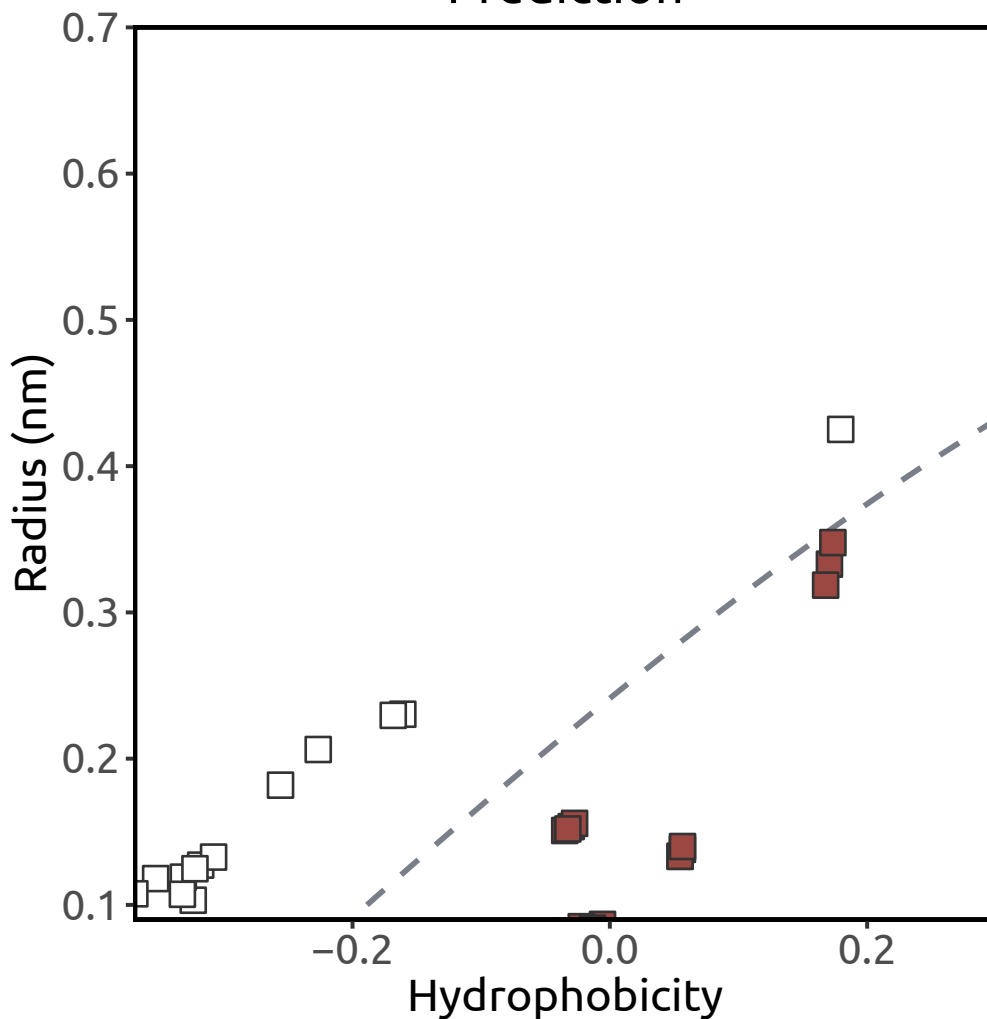
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.63 (n = 19)

Simulation result:
barrier to water

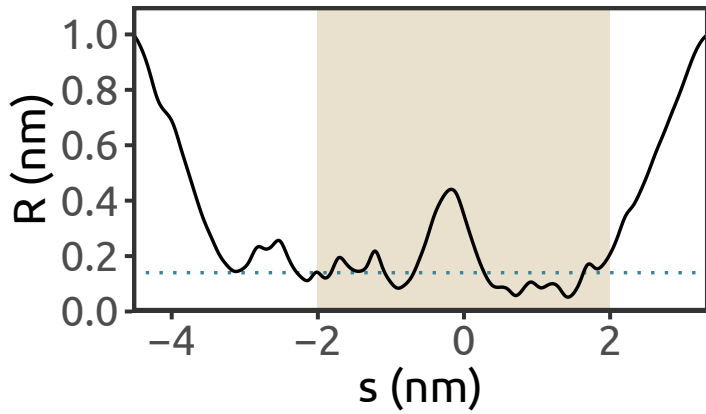
KcsA (PDB ID: 2ITD)

Streptomyces lividans

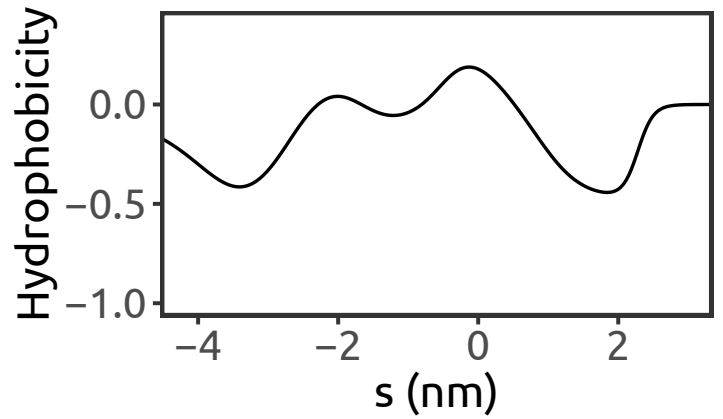
X-ray (2.7 Å)

Lockless et al., 2007

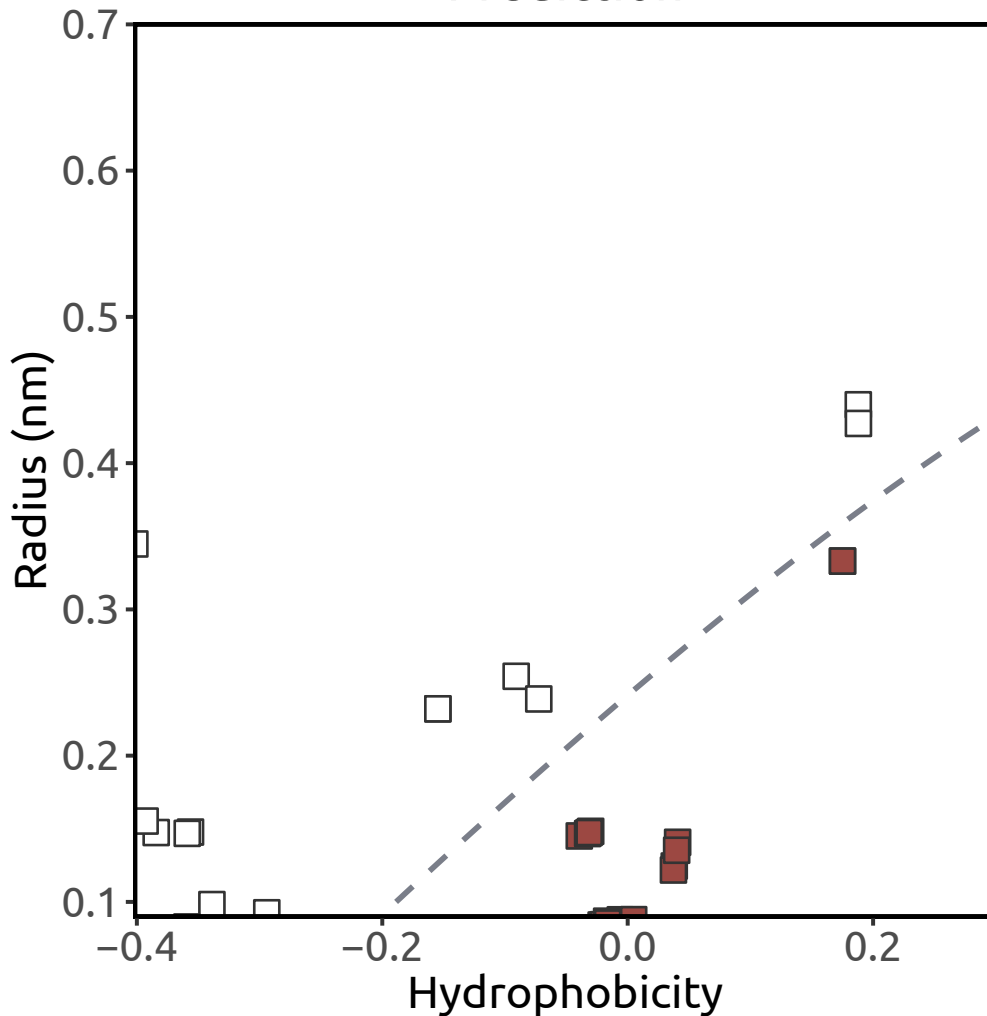
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.65 (n = 18)

Simulation result:

barrier to water

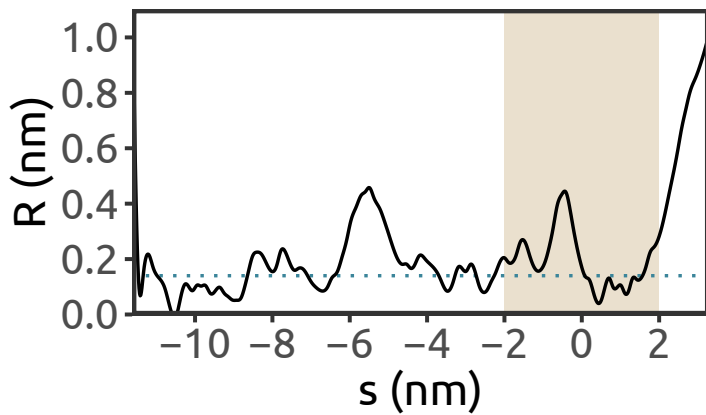
KcsA (PDB ID: 3EFF)

Streptomyces lividans

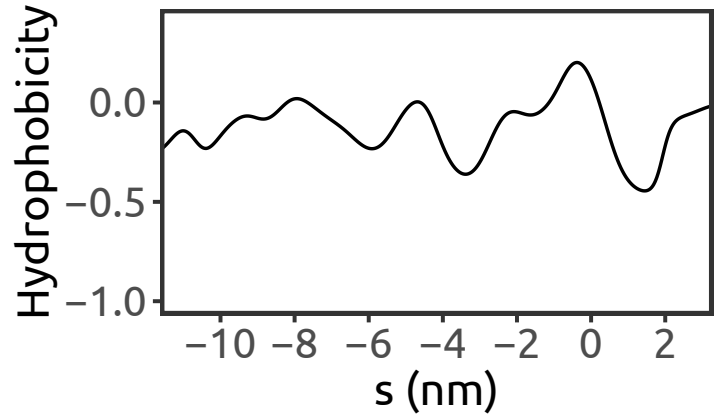
X-ray (3.8 Å)

Uysal et al., 2009

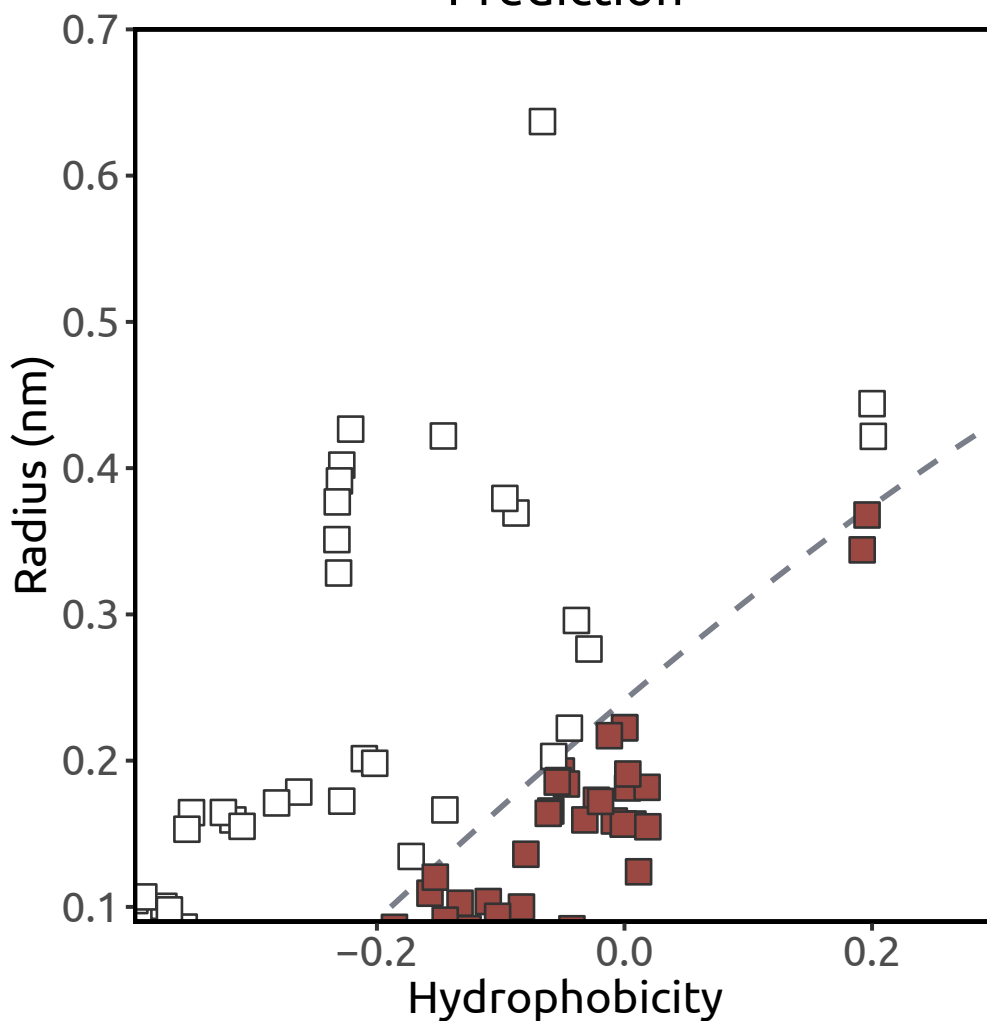
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.95 (n = 41)

Simulation result:
barrier to water

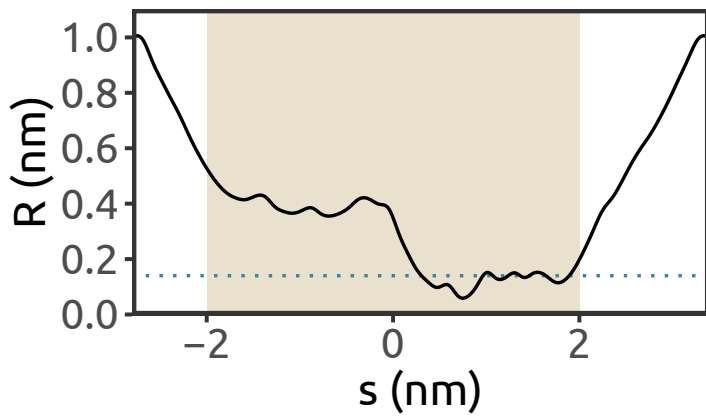
KcsA (PDB ID: 3FB5)

Streptomyces lividans

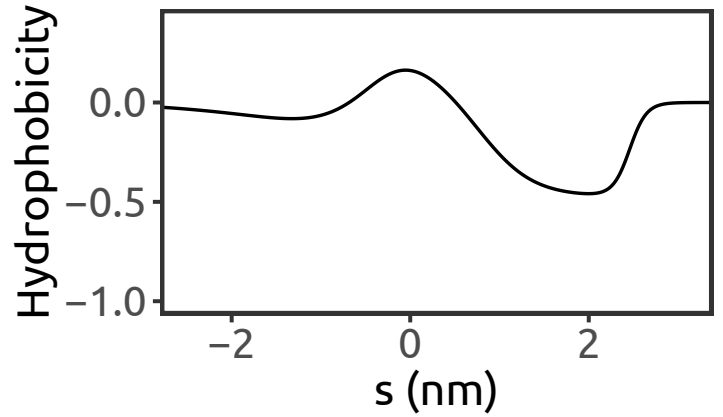
X-ray (2.8 Å)

Cuello et al., 2010

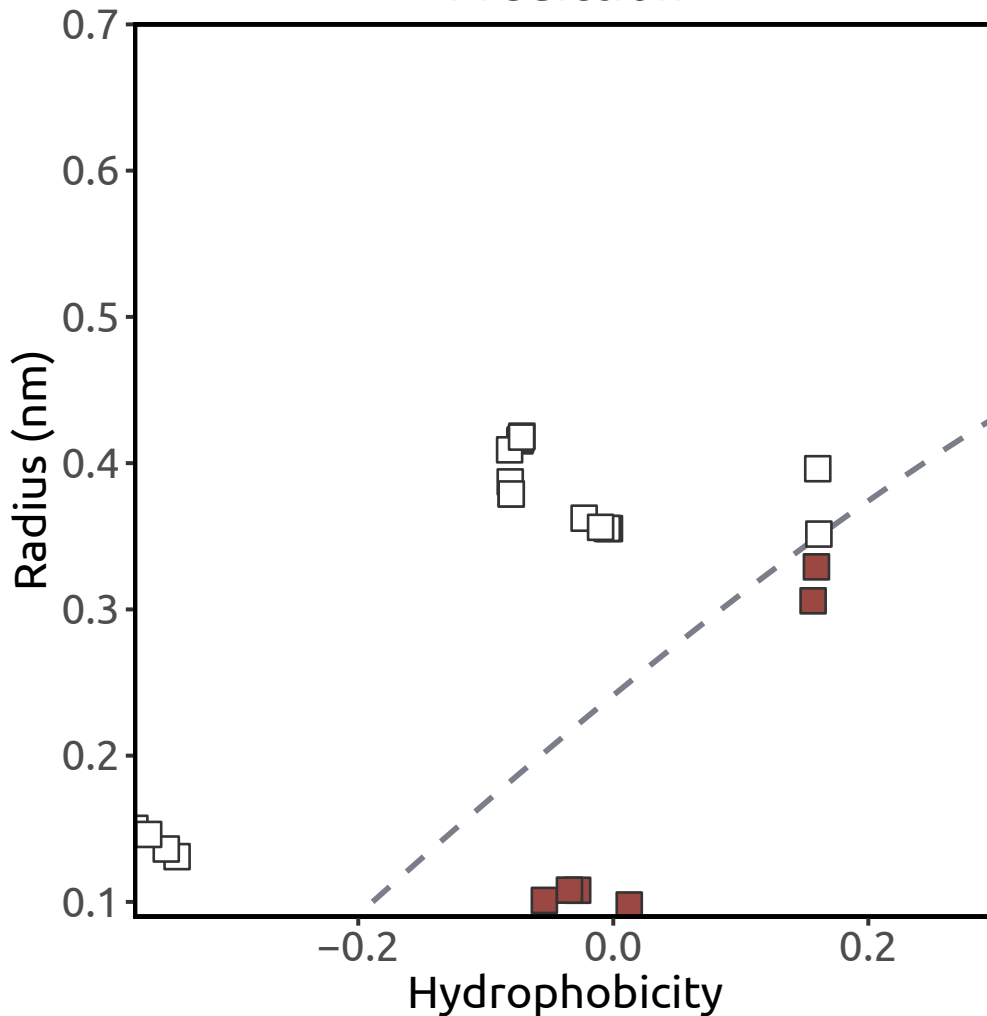
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.43 (n = 6)

Simulation result:
hydrated channel

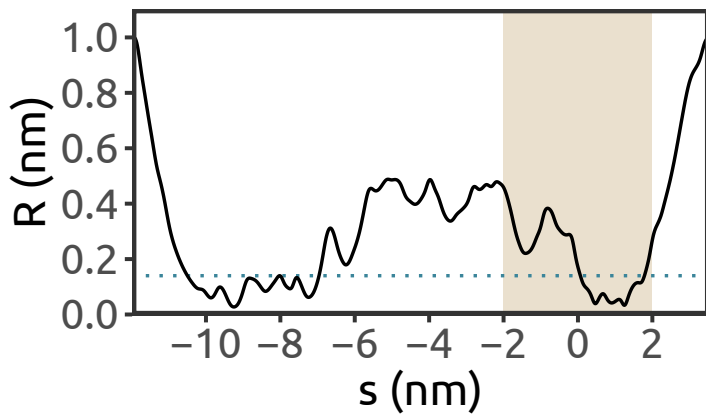
KcsA (PDB ID: 3PJS)

Streptomyces lividans

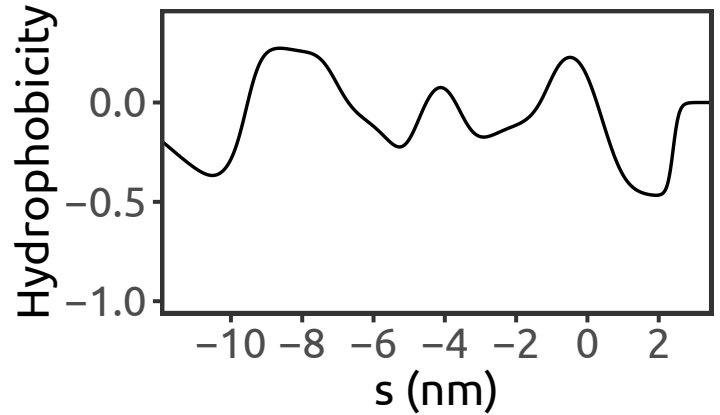
X-ray (3.8 Å)

Uysal et al., 2011

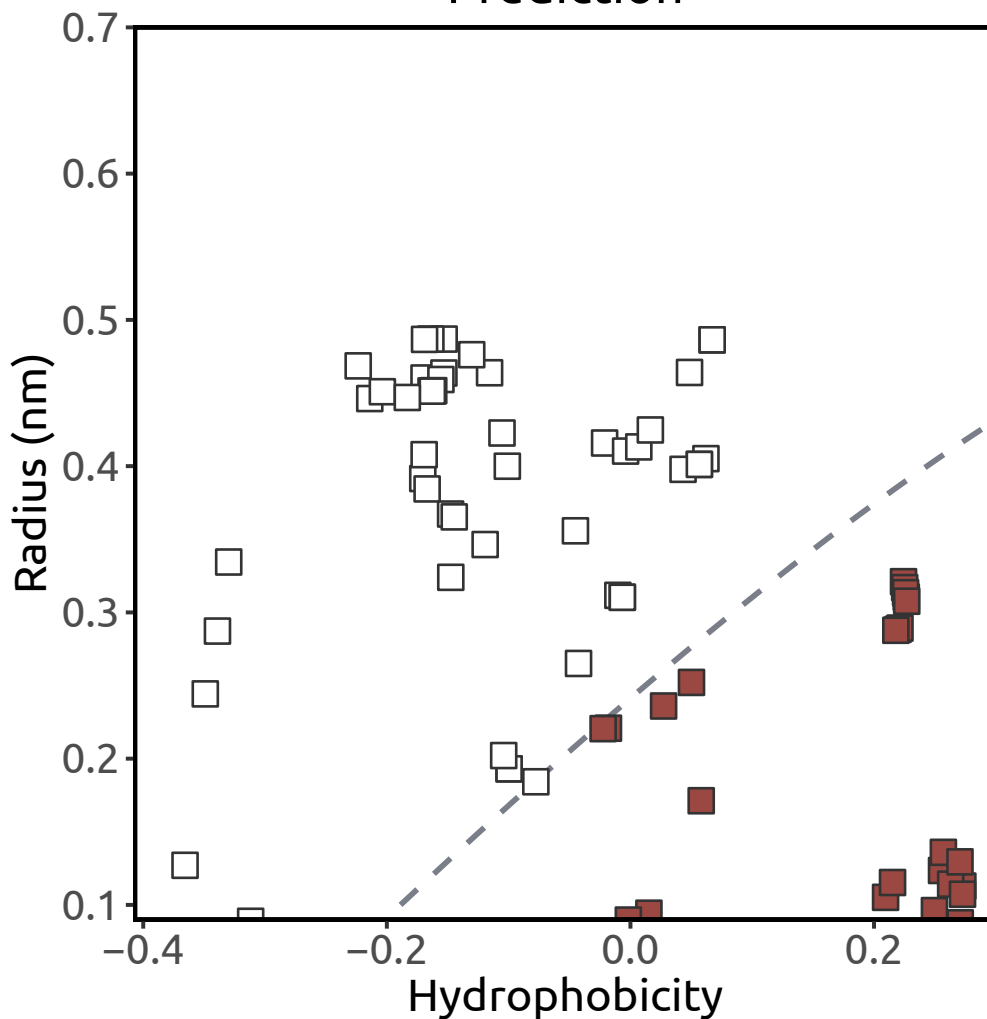
Pore radius



Hydrophobicity



Prediction



Heuristic score:

5 (n = 32)

Simulation result:

barrier to water

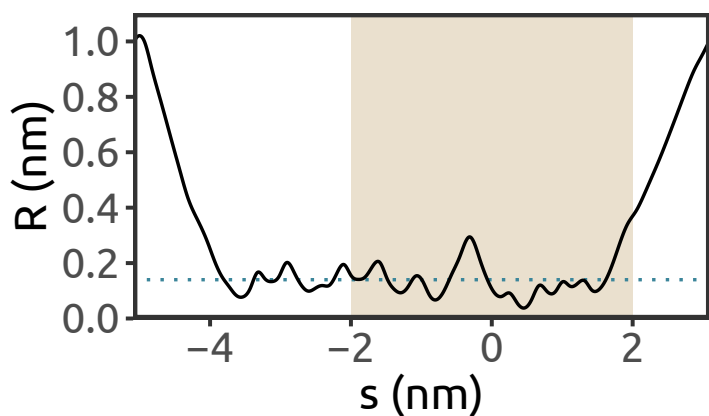
KcsA (PDB ID: 4UUJ)

Streptomyces lividans

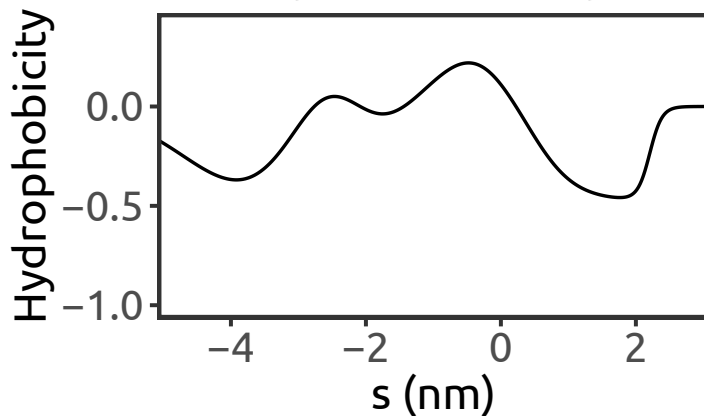
X-ray (2.4 Å)

Lenaeus et al., 2014

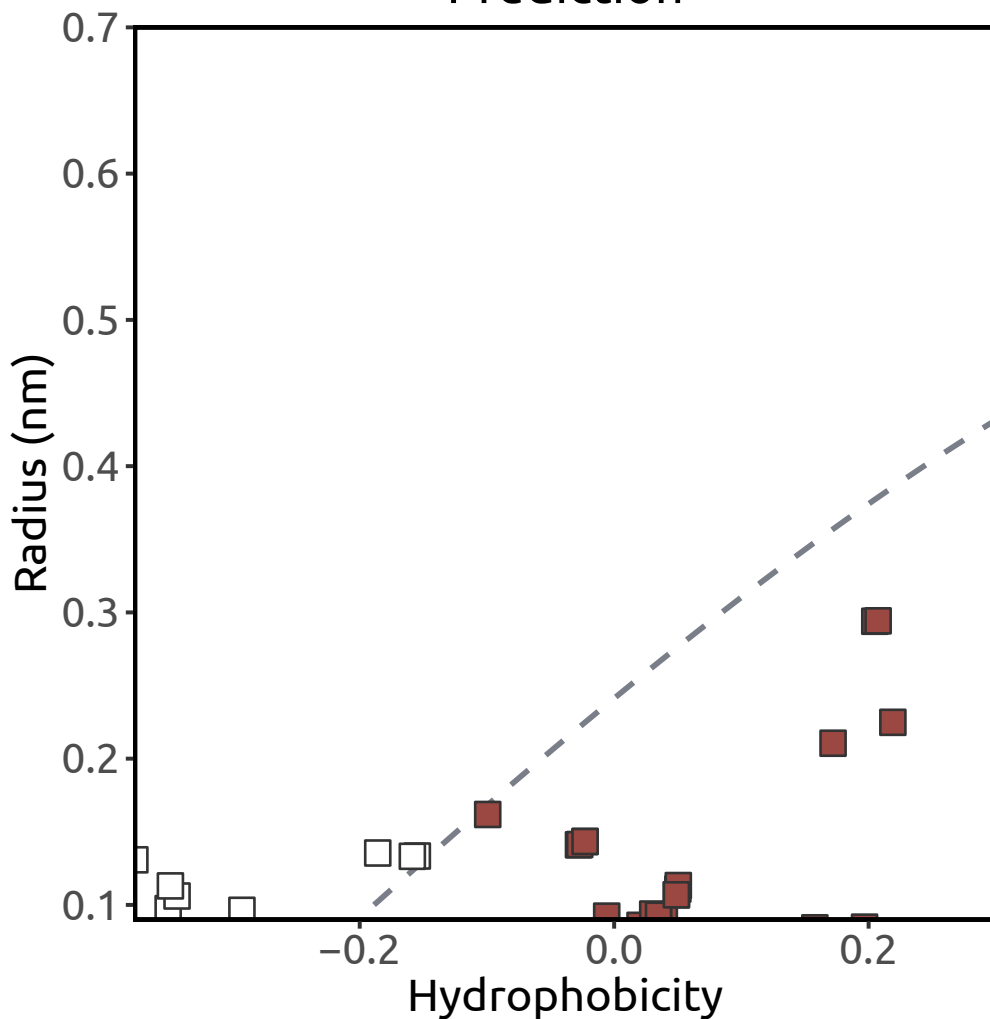
Pore radius



Hydrophobicity



Prediction



Heuristic score:
3.17 (n = 25)

Simulation result:
barrier to water

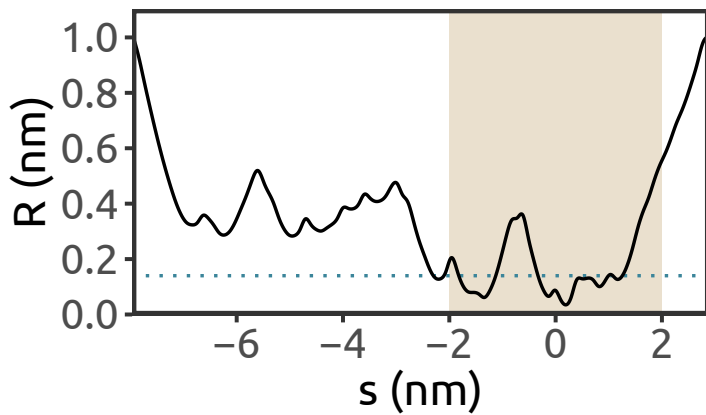
Kir2.2 (PDB ID: 3JYC)

Gallus gallus

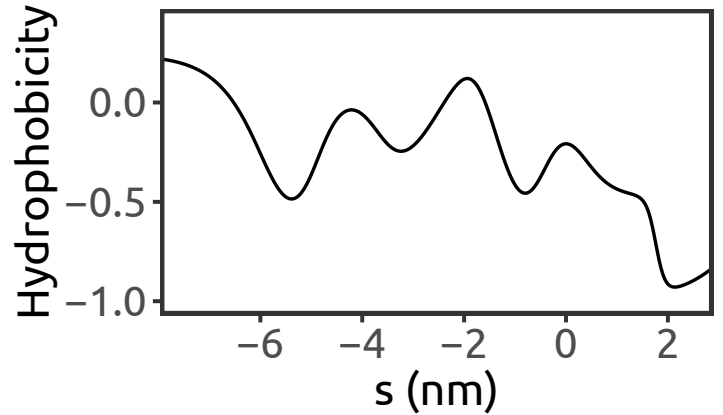
X-ray (3.11 Å)

Tao et al., 2009

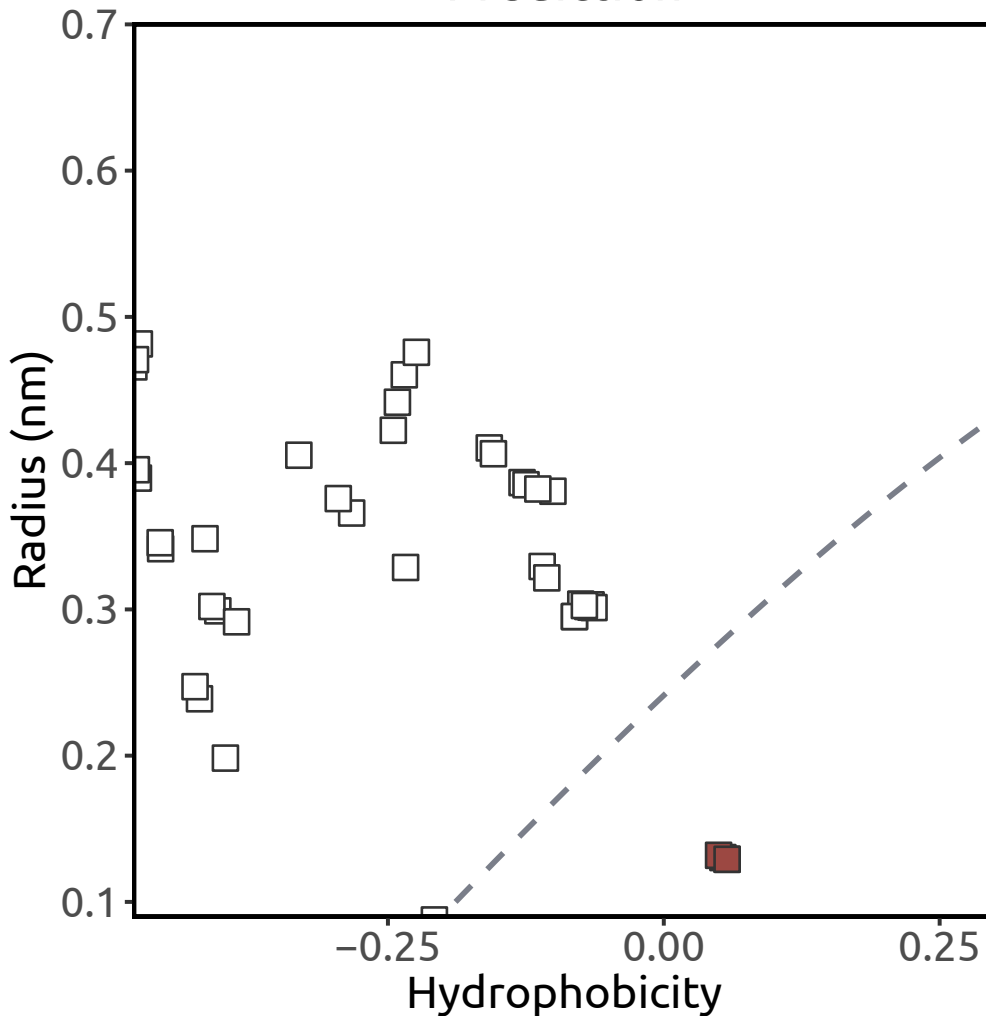
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.88 (n = 8)

Simulation result:

barrier to water

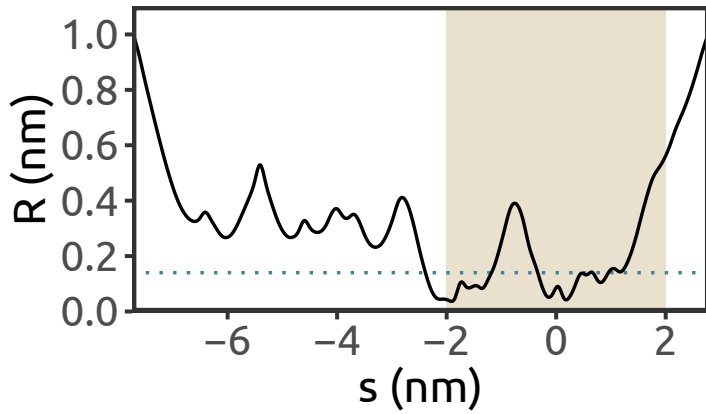
Kir2.2 (PDB ID: 3SPC)

Gallus gallus

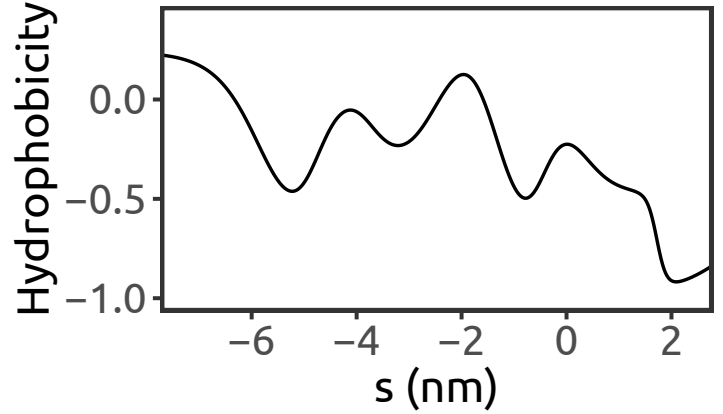
X-ray (2.45 Å)

Hansen et al., 2011

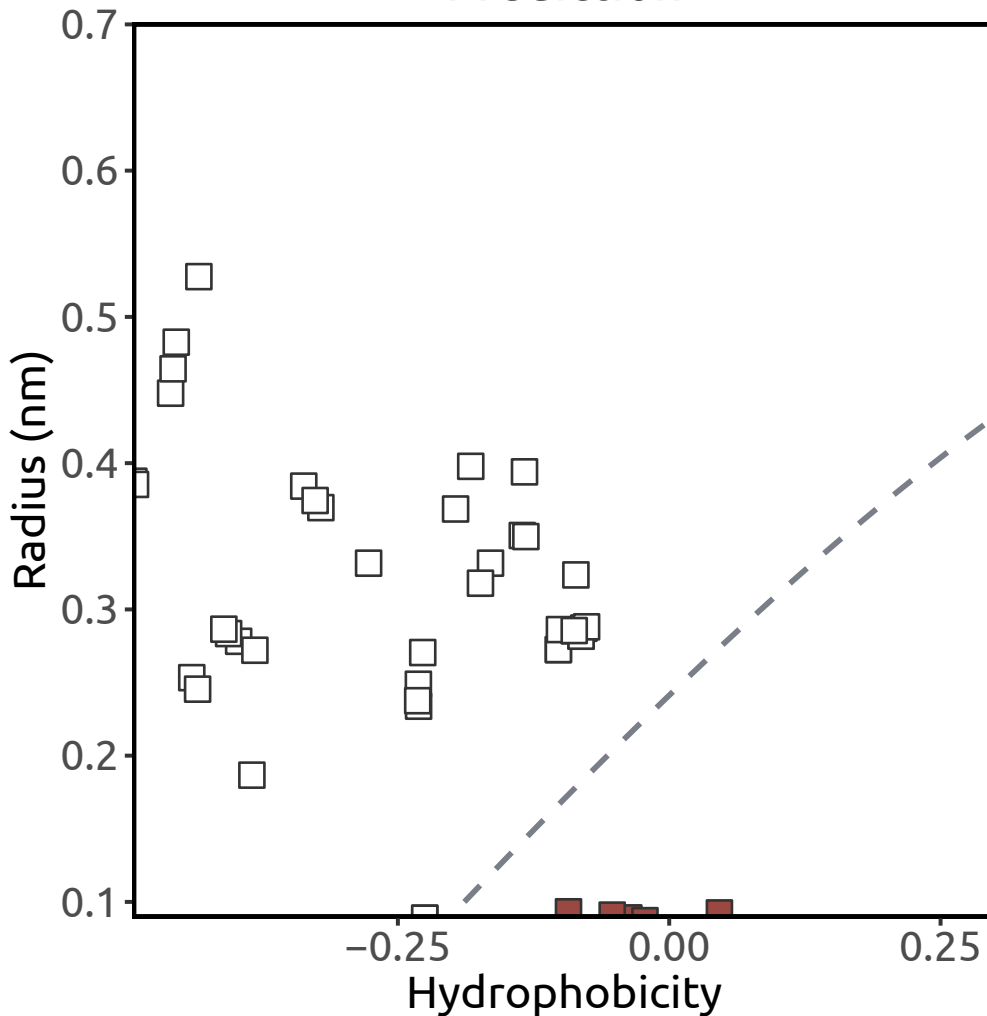
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.15 (n = 8)

Simulation result:

barrier to water

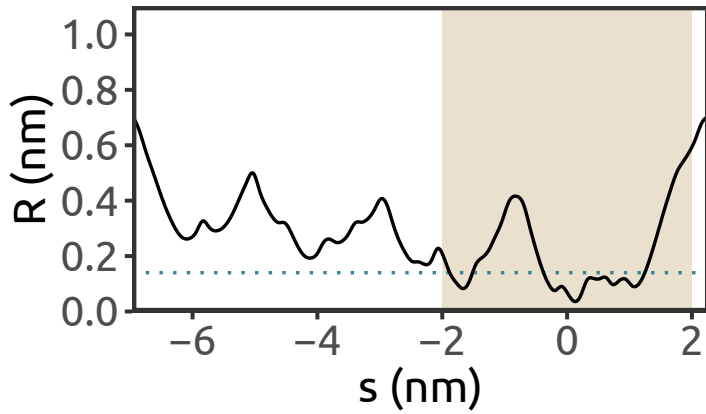
Kir2.2 (PDB ID: 3SPI)

Gallus gallus

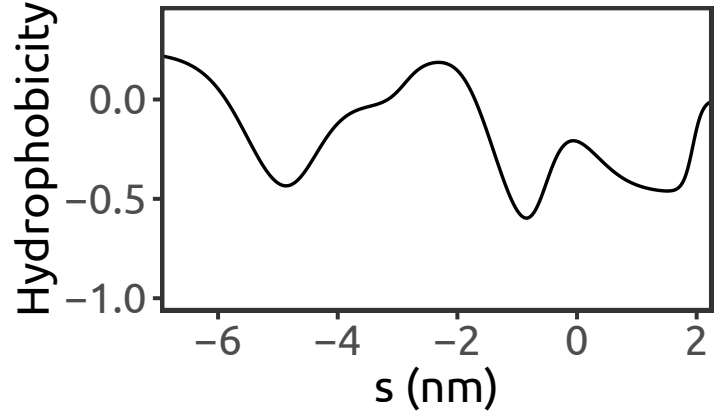
X-ray (3.31 Å)

Hansen et al., 2011

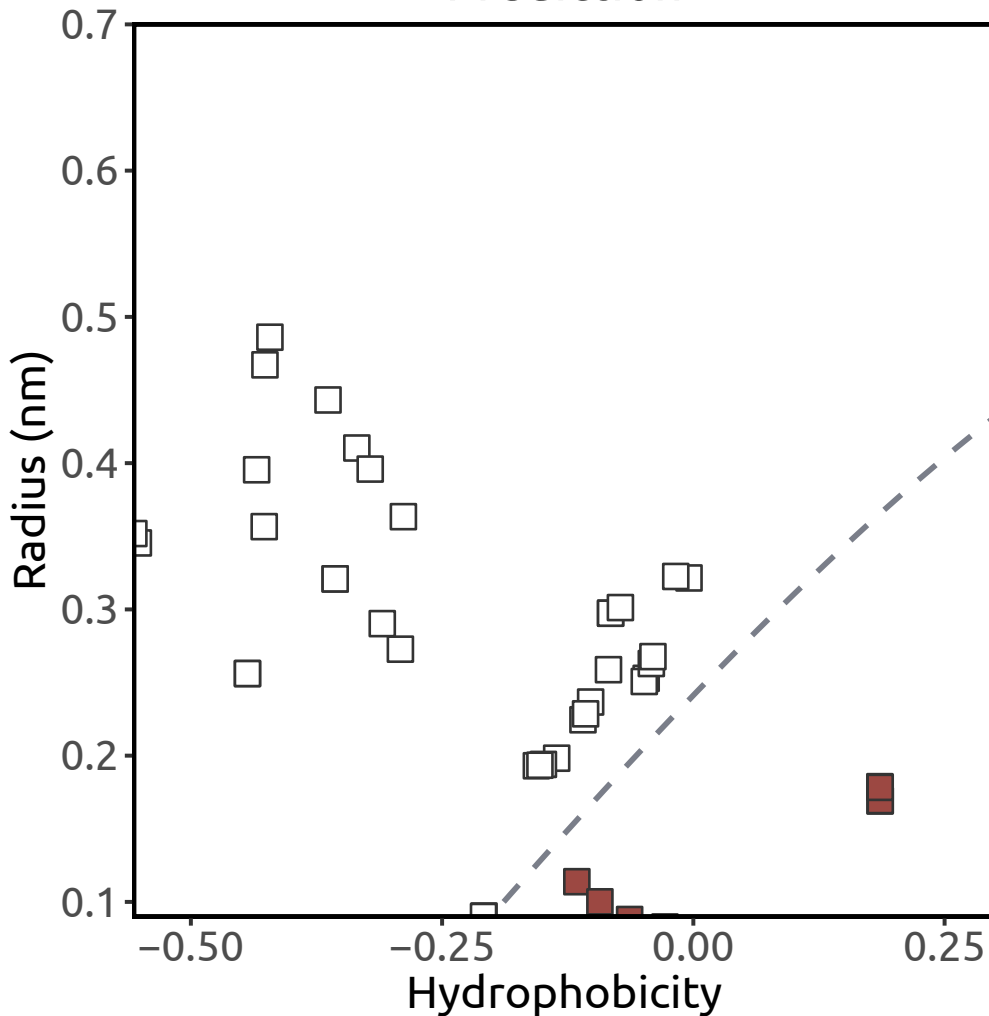
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.93 (n = 8)

Simulation result:

barrier to water

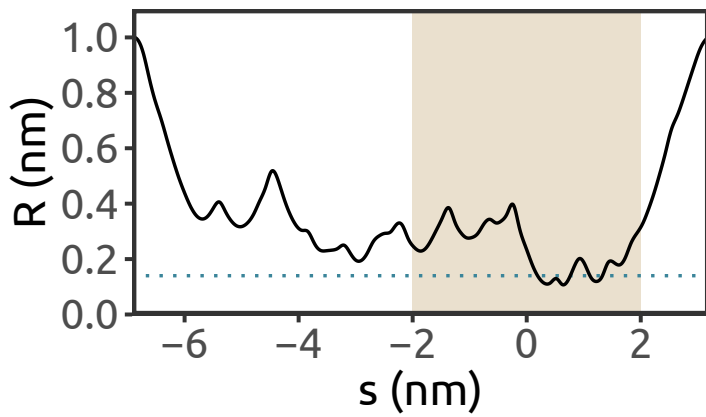
Kir3.2 (PDB ID: 3SYA)

Mus musculus

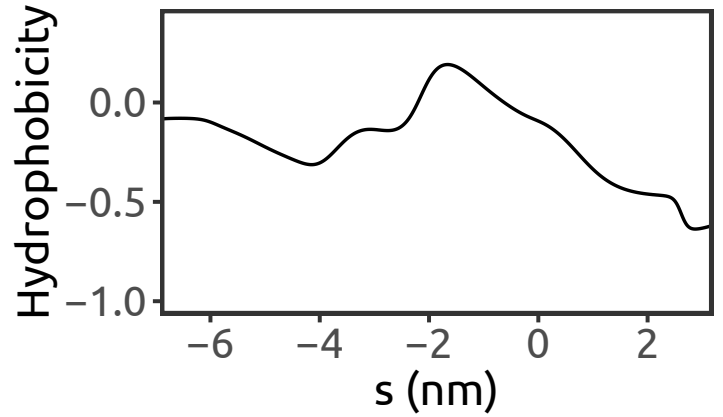
X-ray (2.98 Å)

Whorton & Mackinnon, 2011

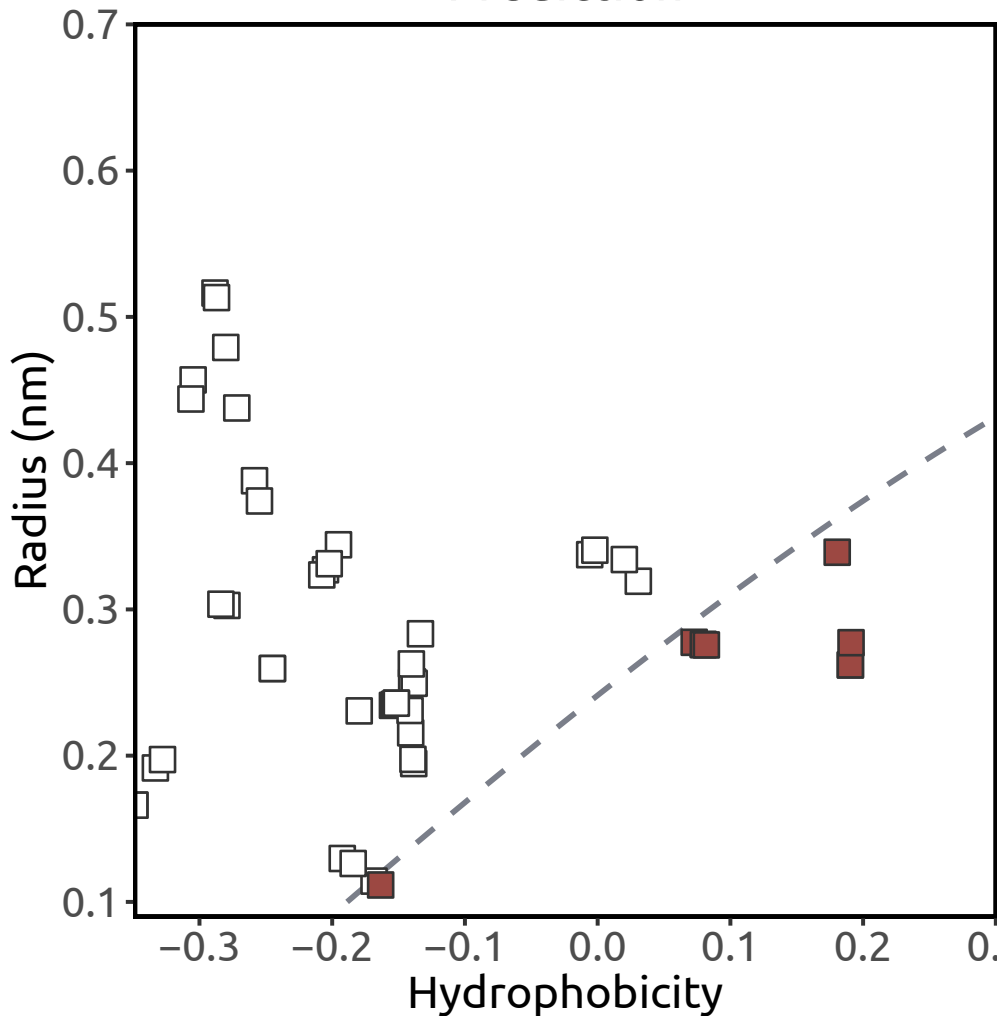
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.35 (n = 9)

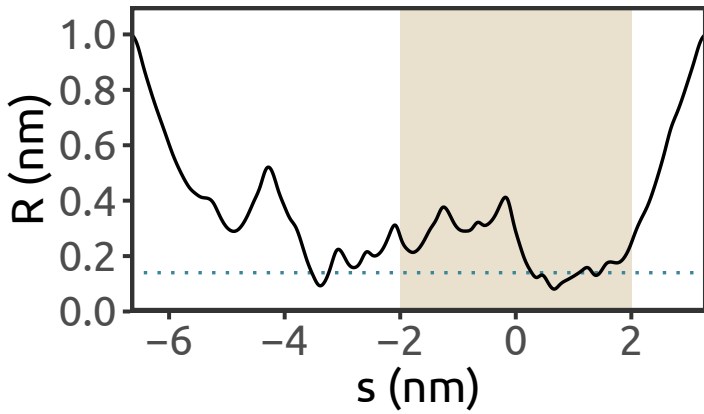
Simulation result:
barrier to water

Kir3.2 (PDB ID: 3SYO)

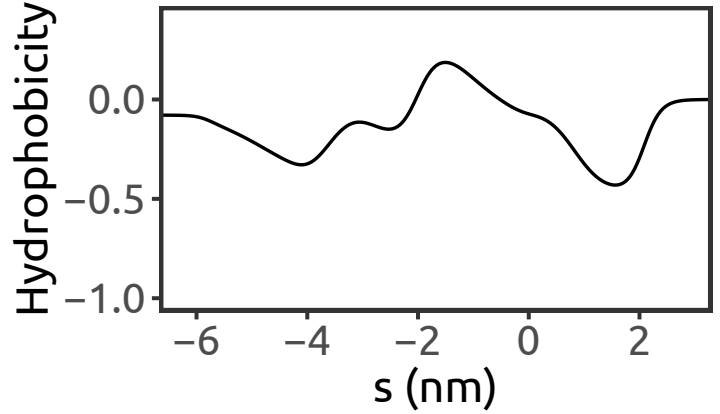
Mus musculus
X-ray (3.54 Å)

Whorton & Mackinnon, 2011

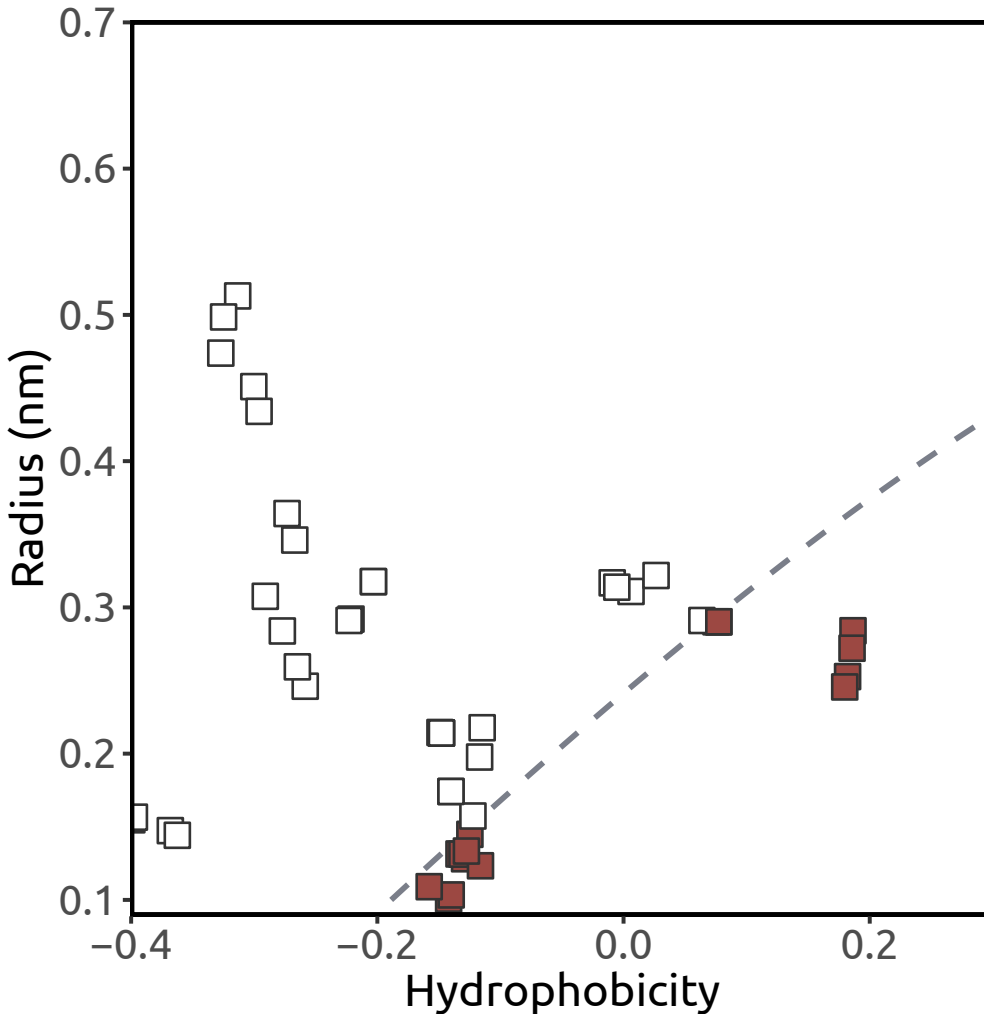
Pore radius



Hydrophobicity



Prediction



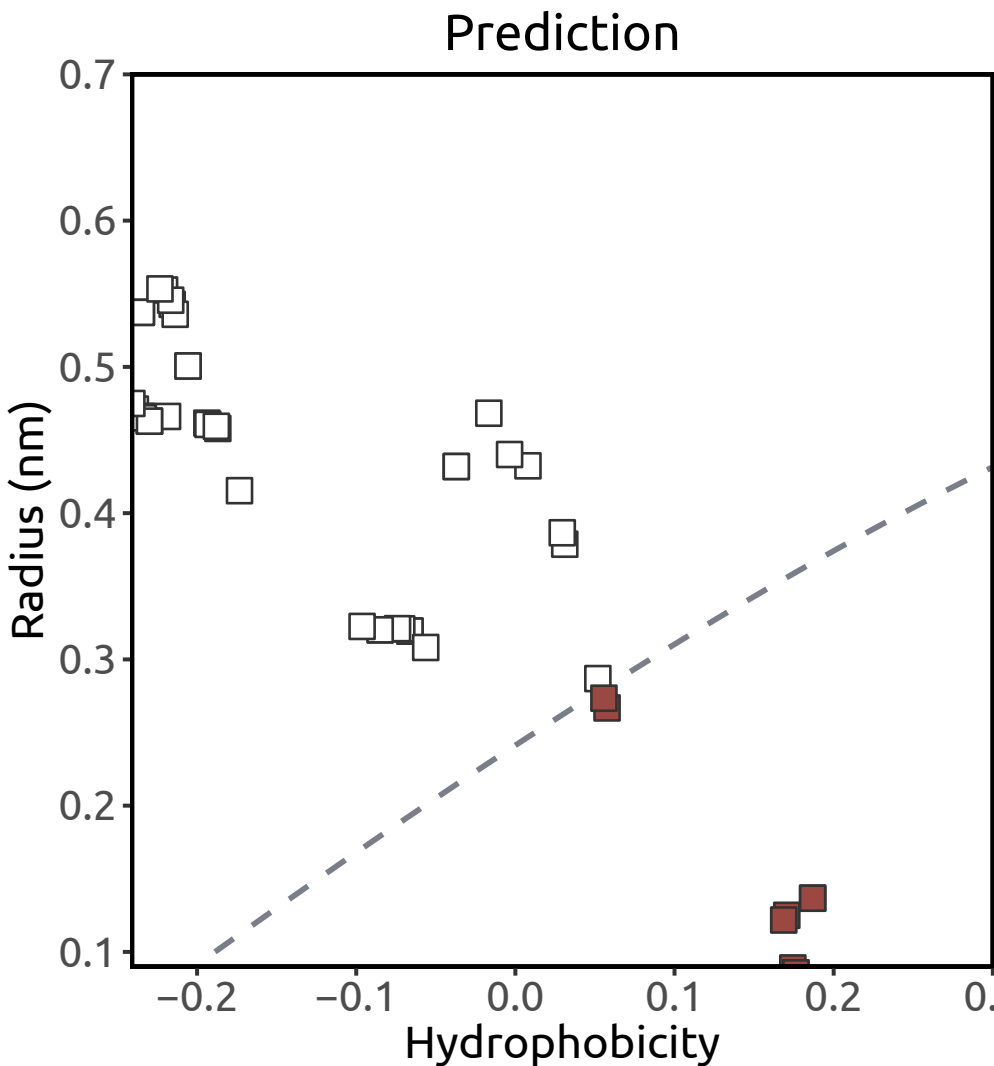
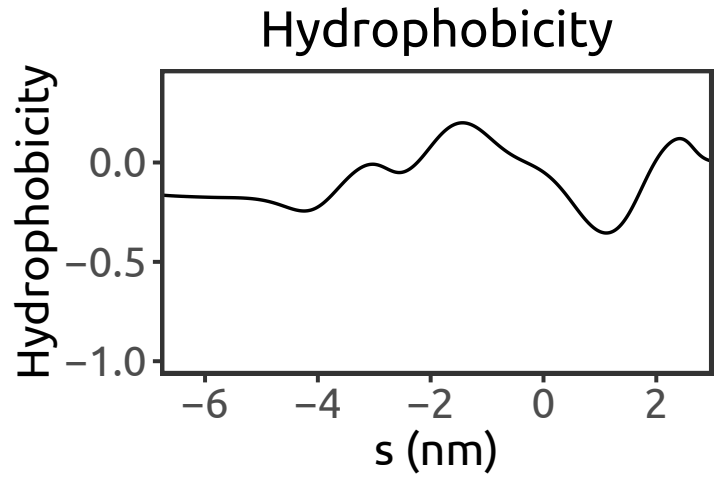
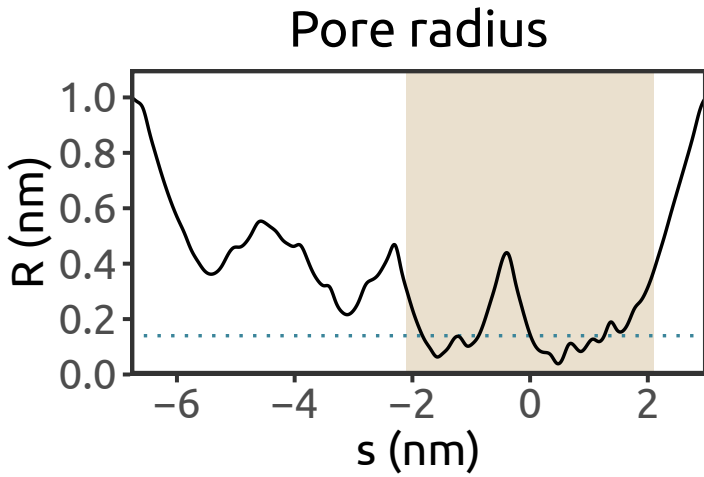
Heuristic score:
0.5 (n = 16)

Simulation result:
hydrated channel

Kir6.2 (PDB ID: 6BAA)

Rattus norvegicus
cryo-EM (3.63 Å)

Martin et al., 2017



Heuristic score:
1.71 (n = 10)

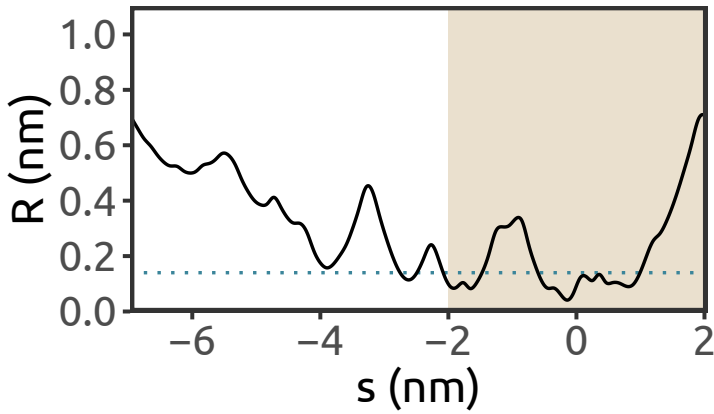
Simulation result:
barrier to water

Kir6.2 (PDB ID: 6C3O)

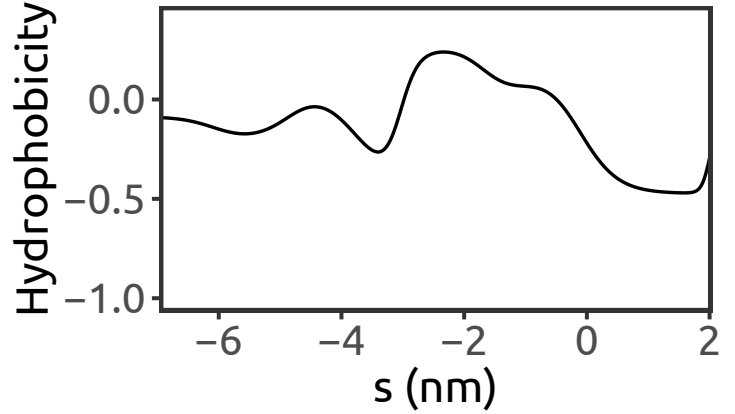
Homo sapiens
cryo-EM (3.9 Å)

Lee et al., 2017

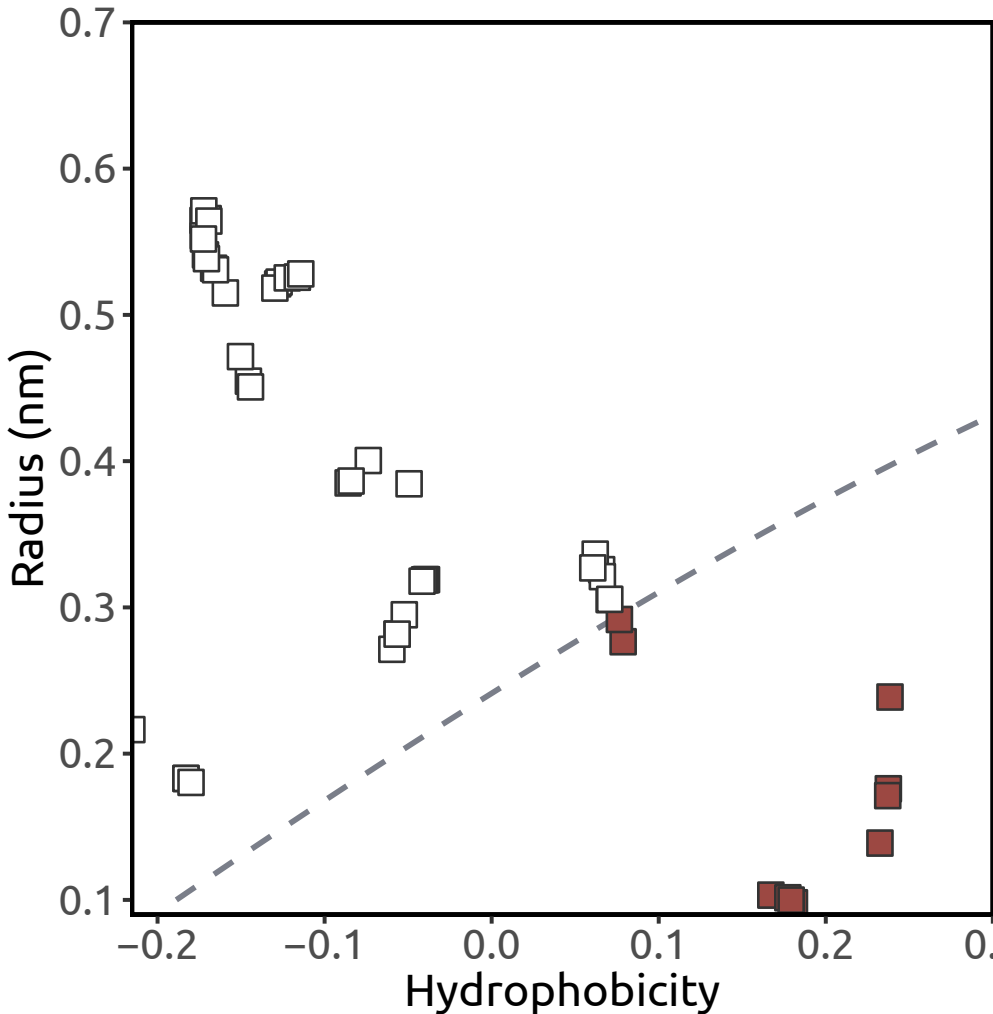
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.62 (n = 10)

Simulation result:
barrier to water

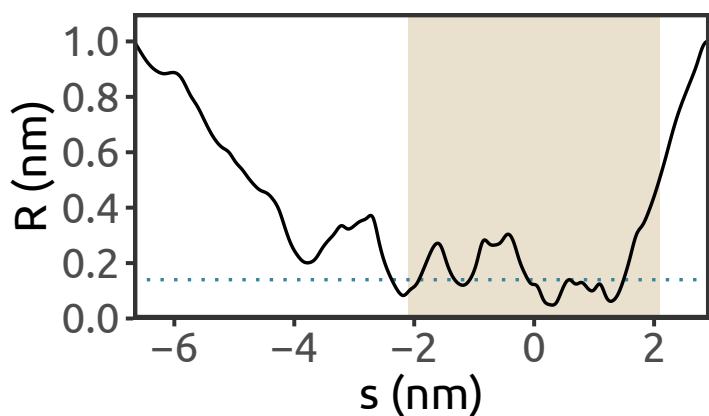
KirBac1.1 (PDB ID: 2WLL)

Burkholderia pseudomallei

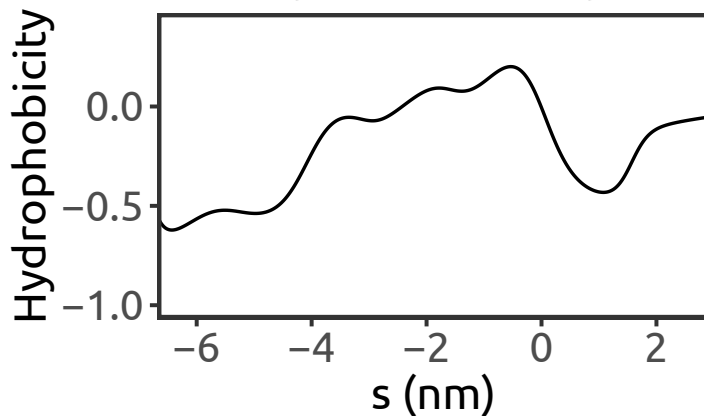
X-ray (3.65 Å)

Kuo et al., 2003

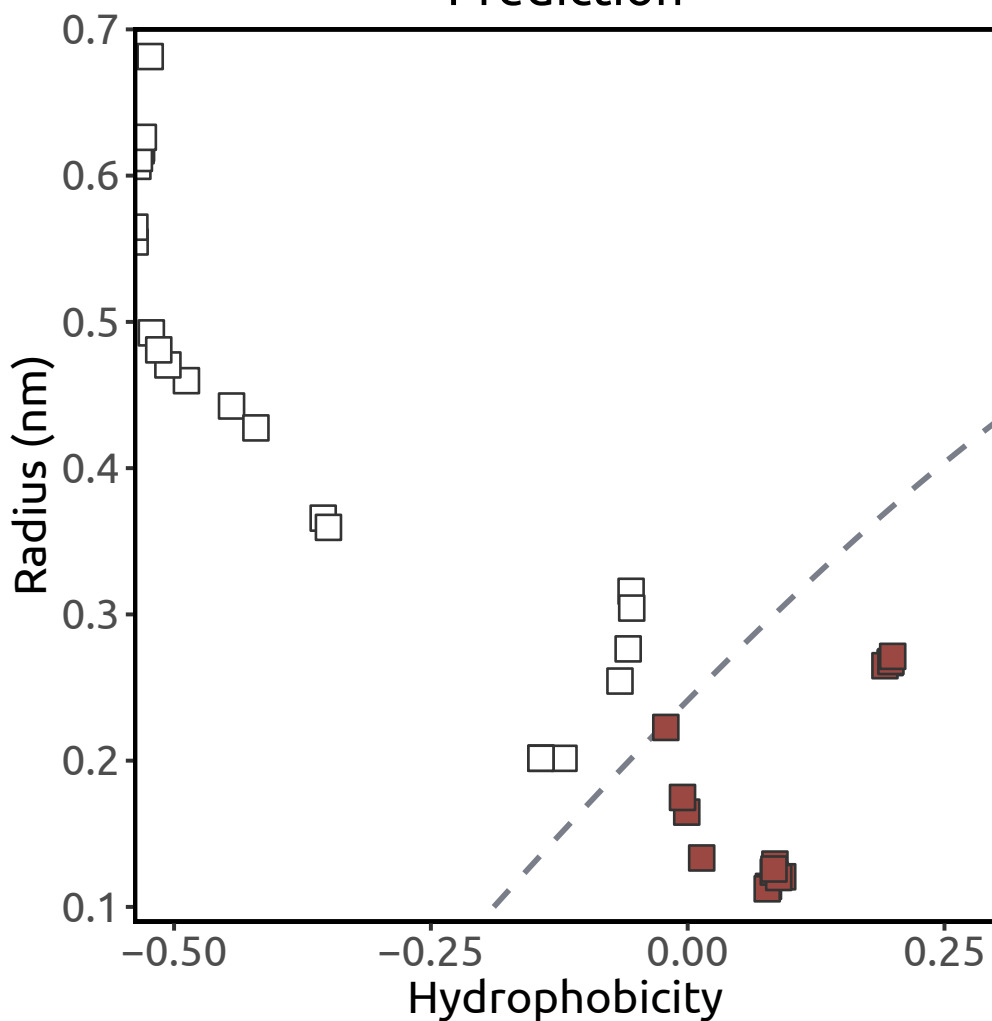
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.74 ($n = 16$)

Simulation result:

barrier to water

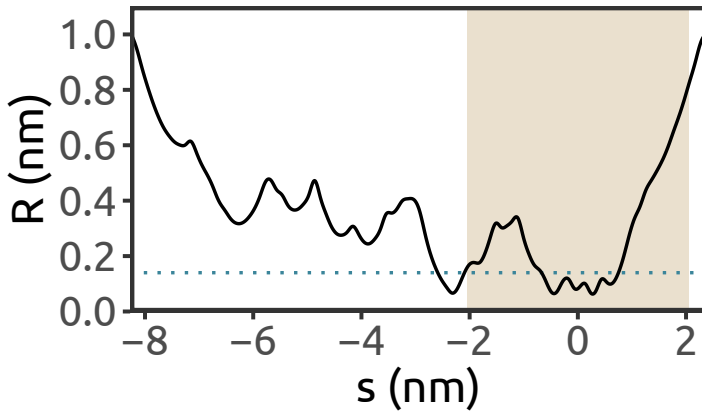
KirBac1.3 (PDB ID: 2QKS)

M. musculus B. xenovornas

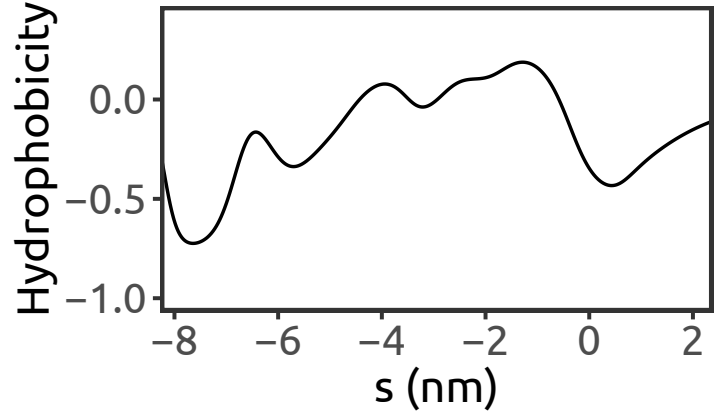
X-ray (2.2 Å)

Nishida et al., 2007

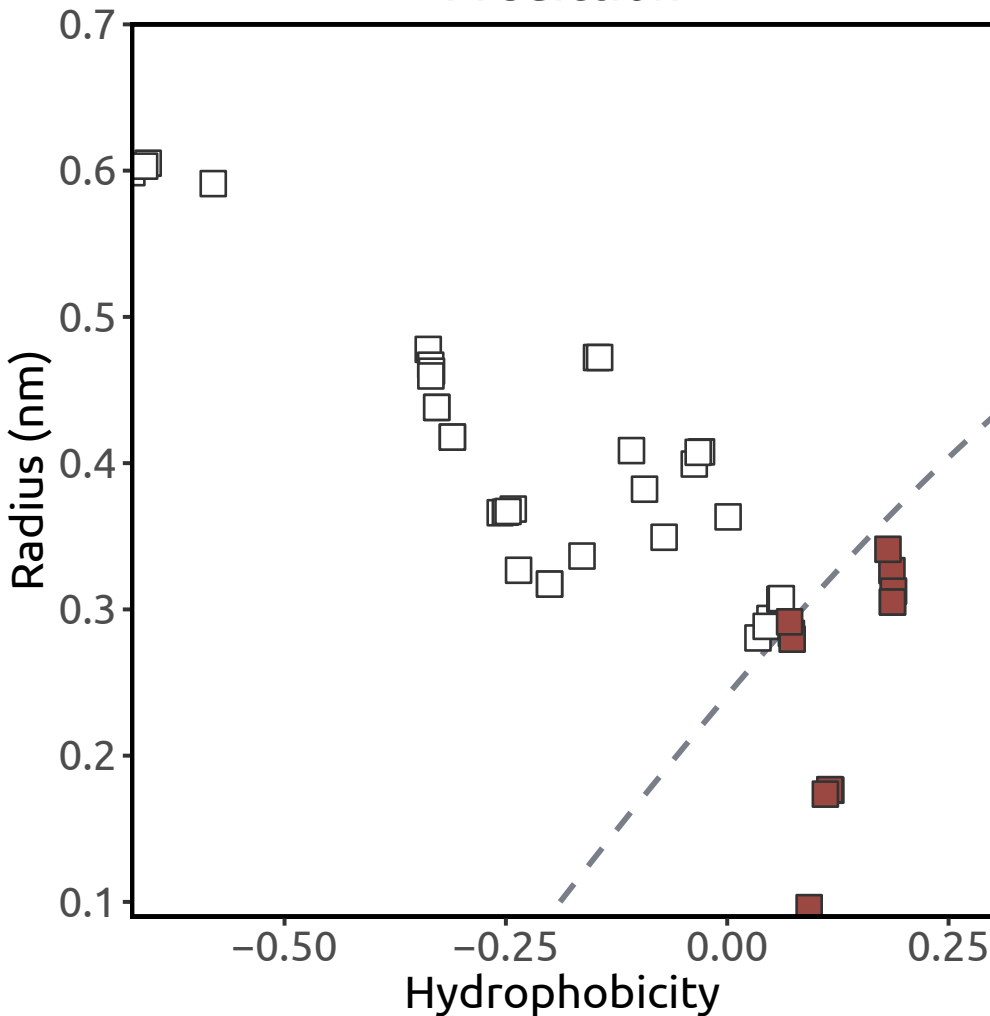
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.49 (n = 17)

Simulation result:

barrier to water

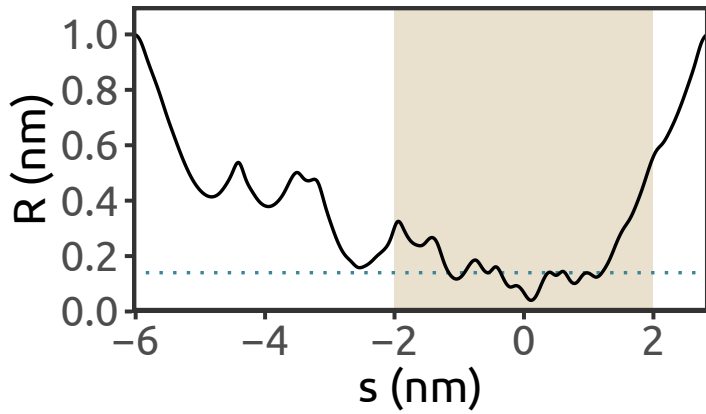
KirBac3.1 (PDB ID: 2WLH)

M. magnetotacticum

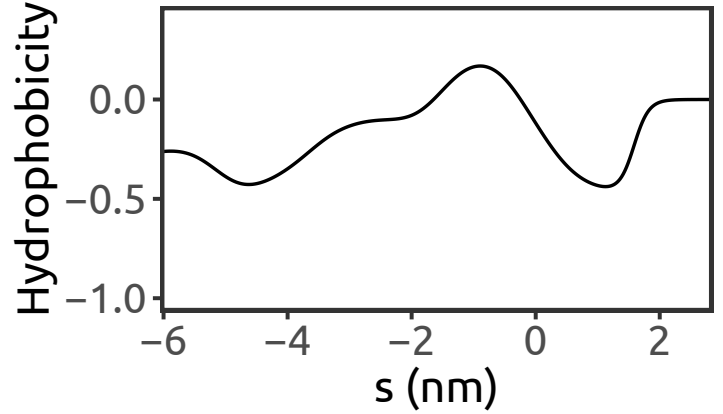
X-ray (3.28 Å)

Clarke et al., 2010

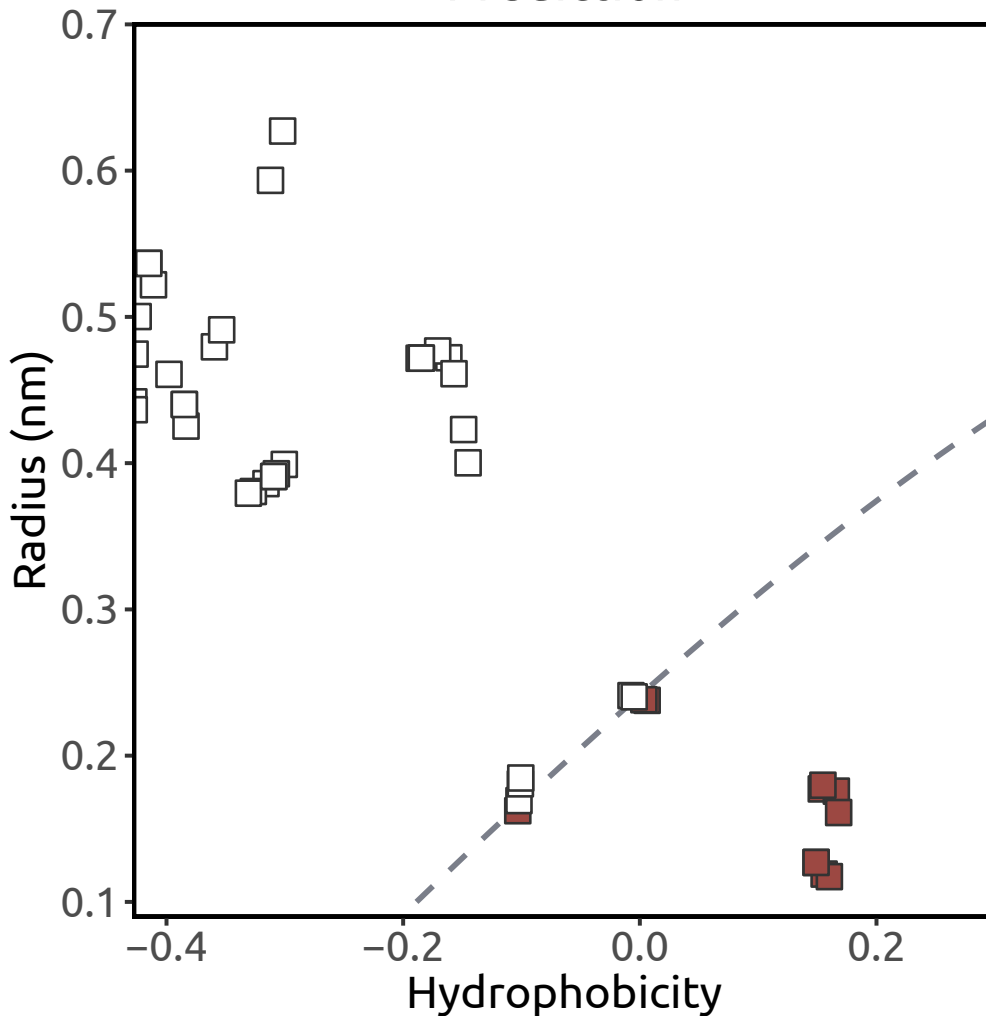
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.35 (n = 11)

Simulation result:

barrier to water

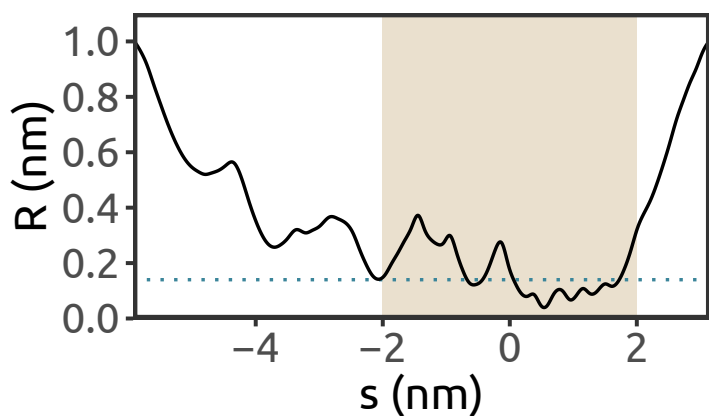
KirBac3.1 (PDB ID: 2WLI)

M. magnetotacticum

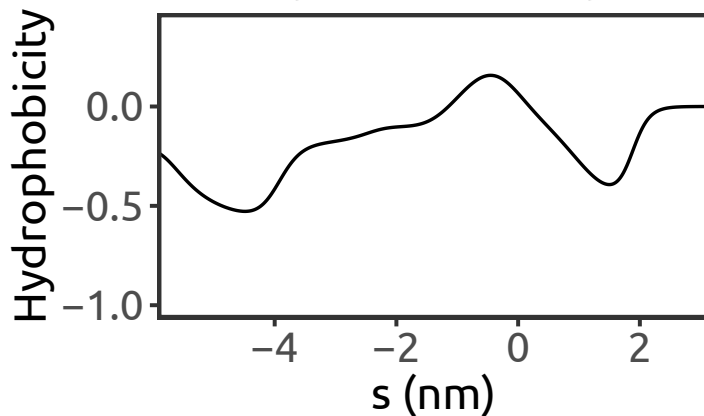
X-ray (3.09 Å)

Clarke et al., 2010

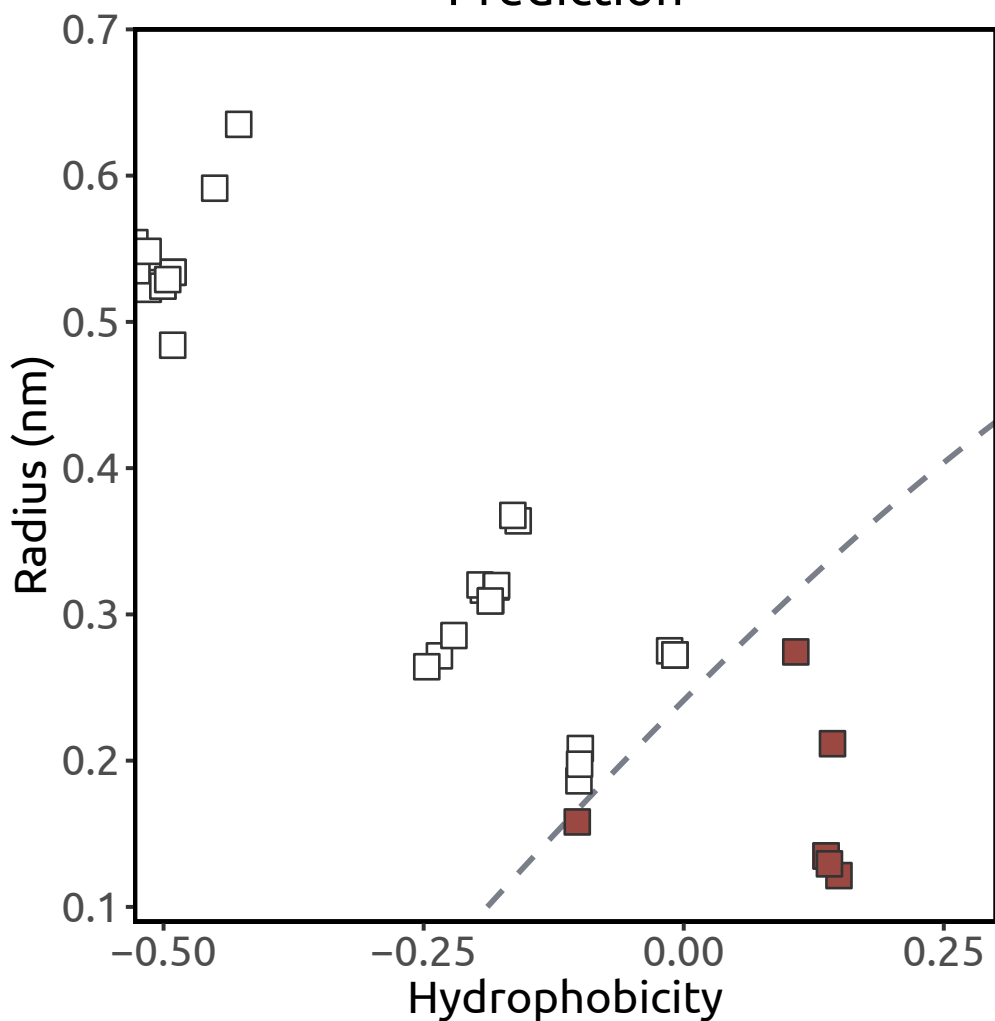
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.84 ($n = 7$)

Simulation result:

barrier to water

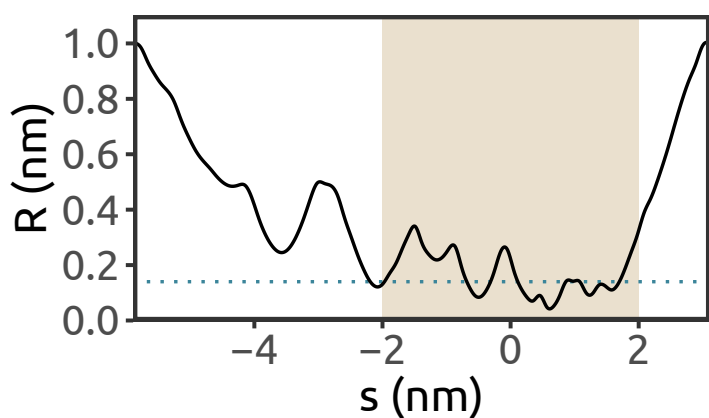
KirBac3.1 (PDB ID: 2WLJ)

M. magnetotacticum

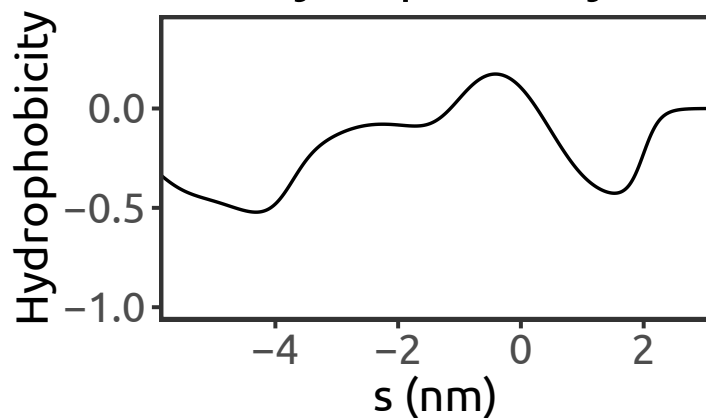
X-ray (2.6 Å)

Clarke et al., 2010

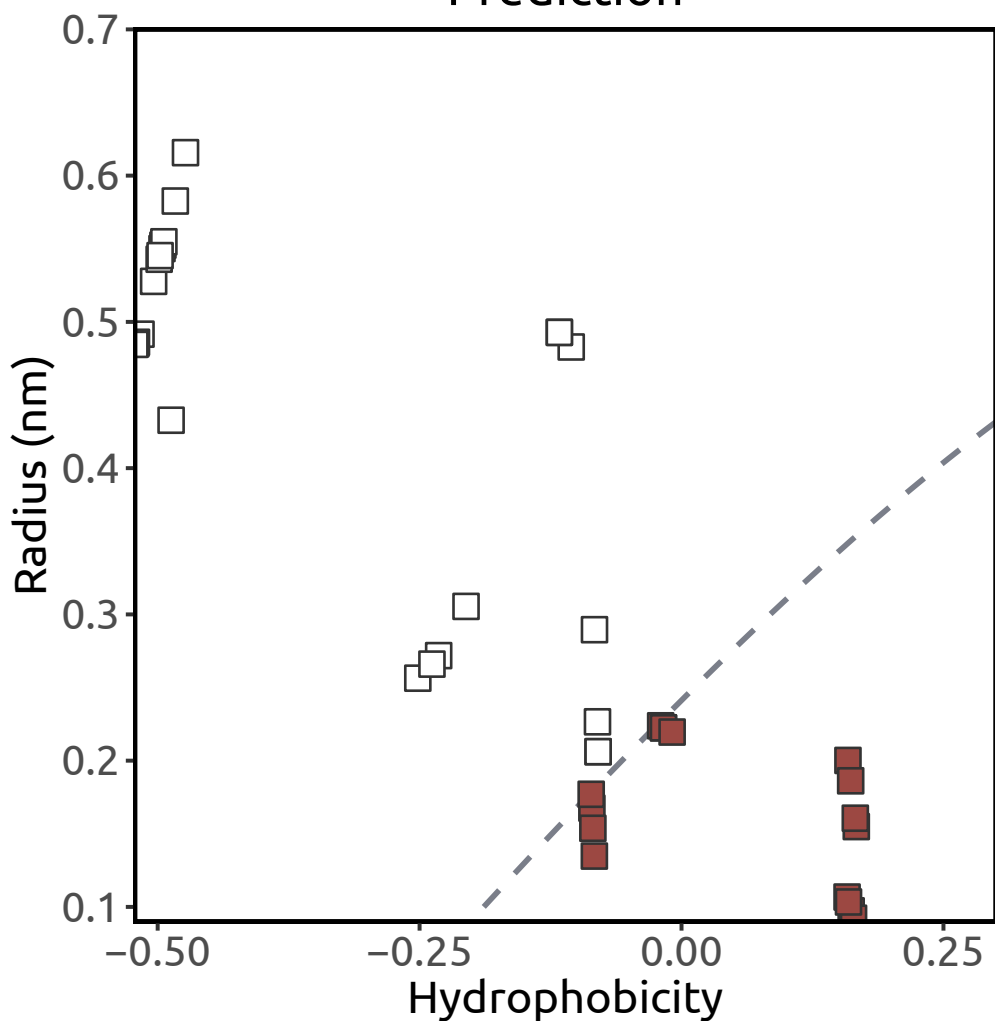
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.52 ($n = 16$)

Simulation result:

barrier to water

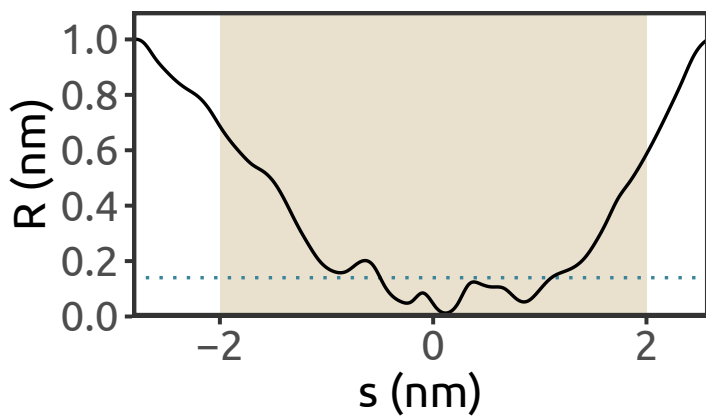
MthK (PDB ID: 1LNQ)

M. thermautotrophicus

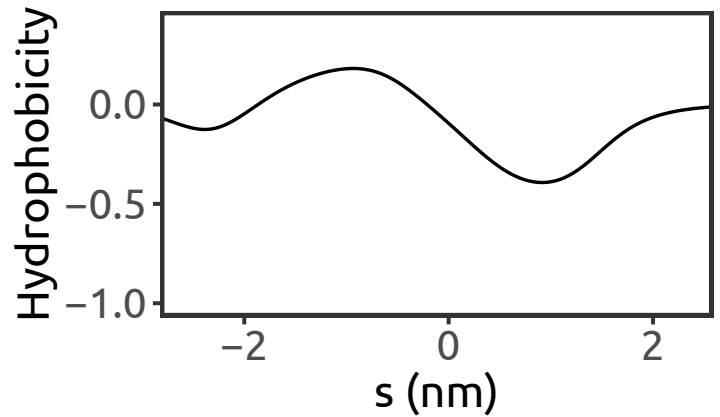
X-ray (3.3 Å)

Jiang et al., 2002

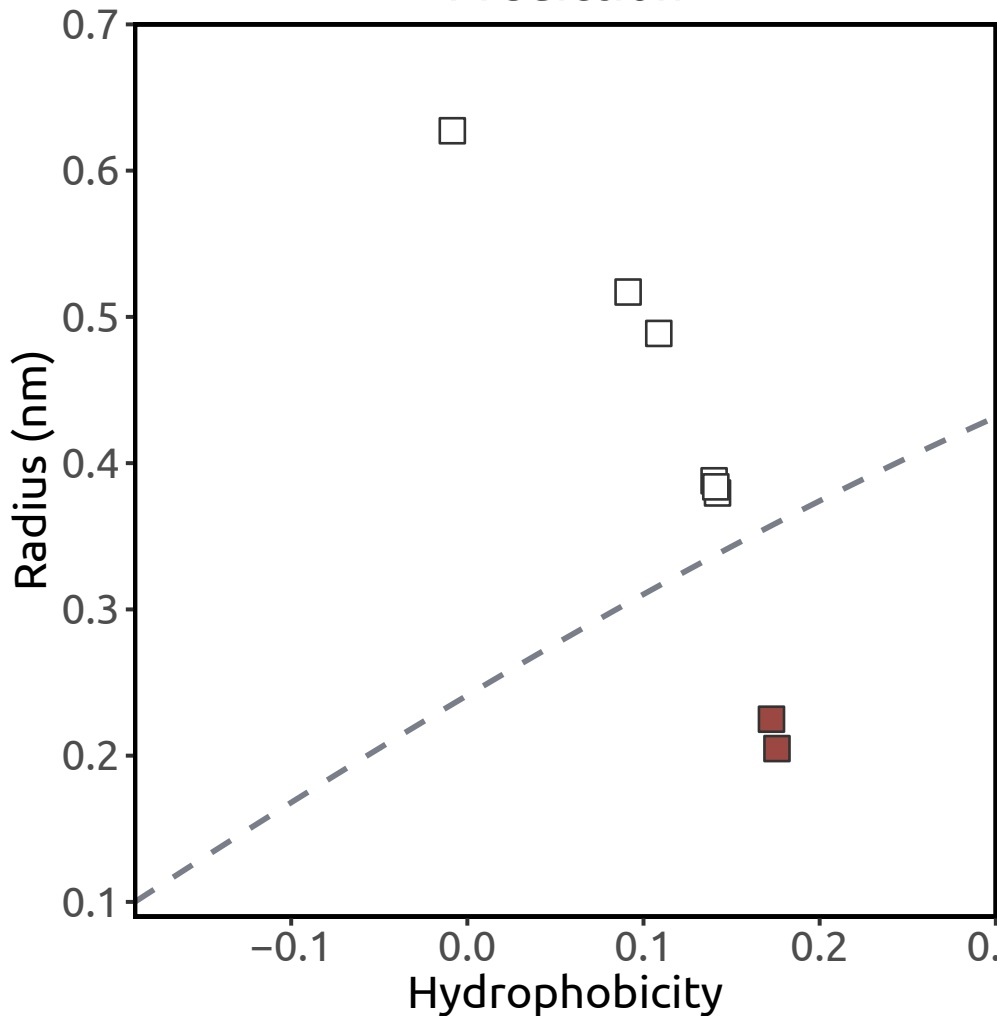
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.24 ($n = 2$)

Simulation result:
hydrated channel

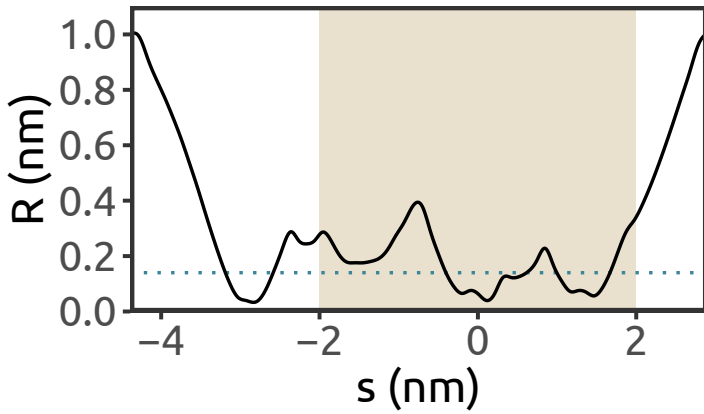
NaK (PDB ID: 2AHY)

Bacillus cereus

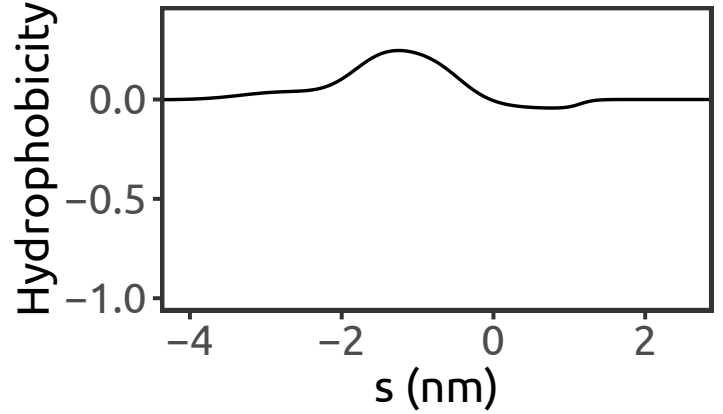
X-ray (2.4 Å)

Shi et al., 2006

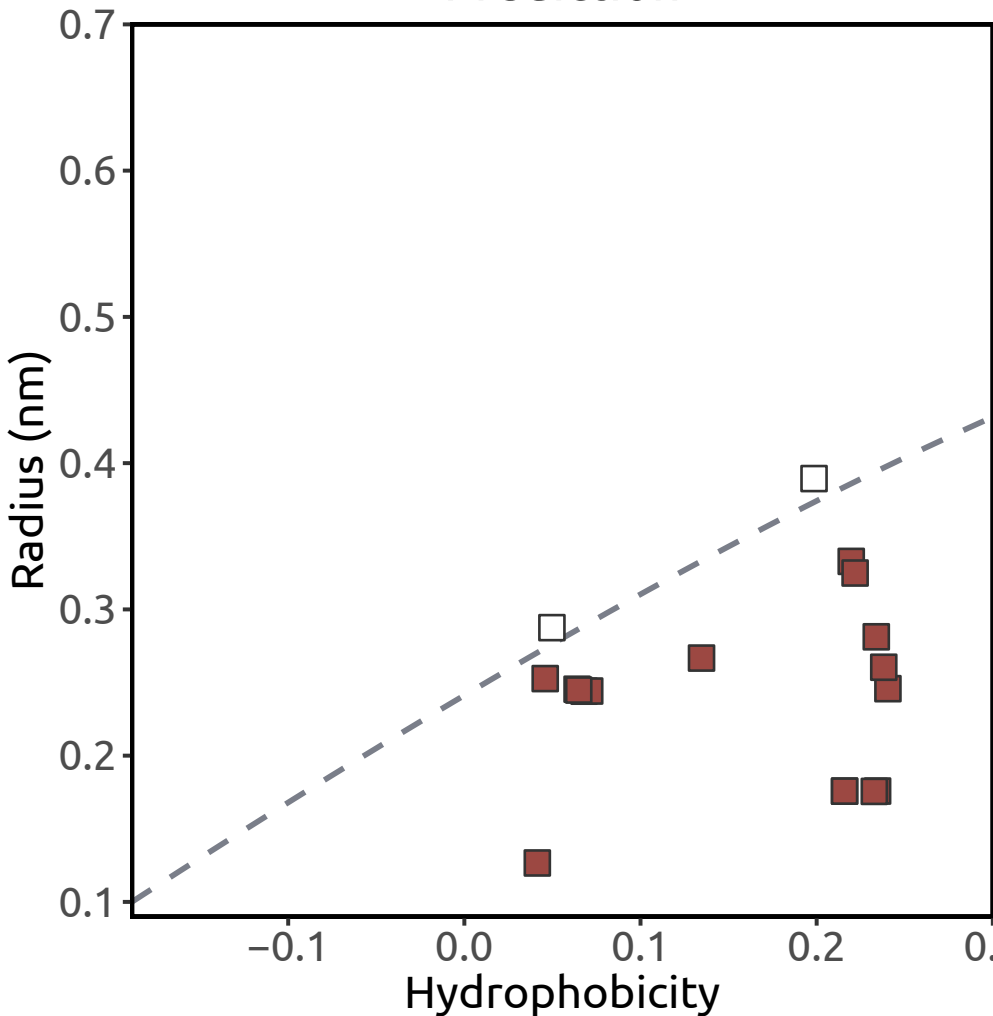
Pore radius



Hydrophobicity



Prediction



Heuristic score:

2.24 (n = 20)

Simulation result:

barrier to water

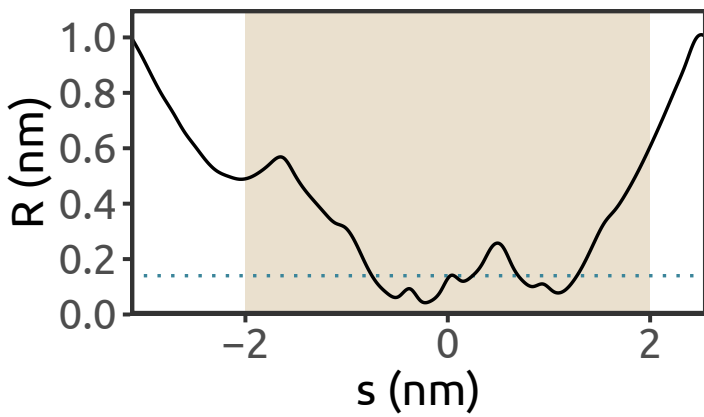
NaK (PDB ID: 3E86)

Bacillus cereus

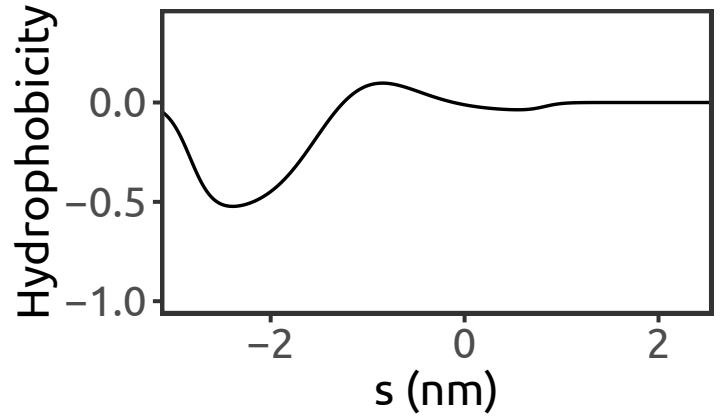
X-ray (1.6 Å)

Alam & Jiang, 2009

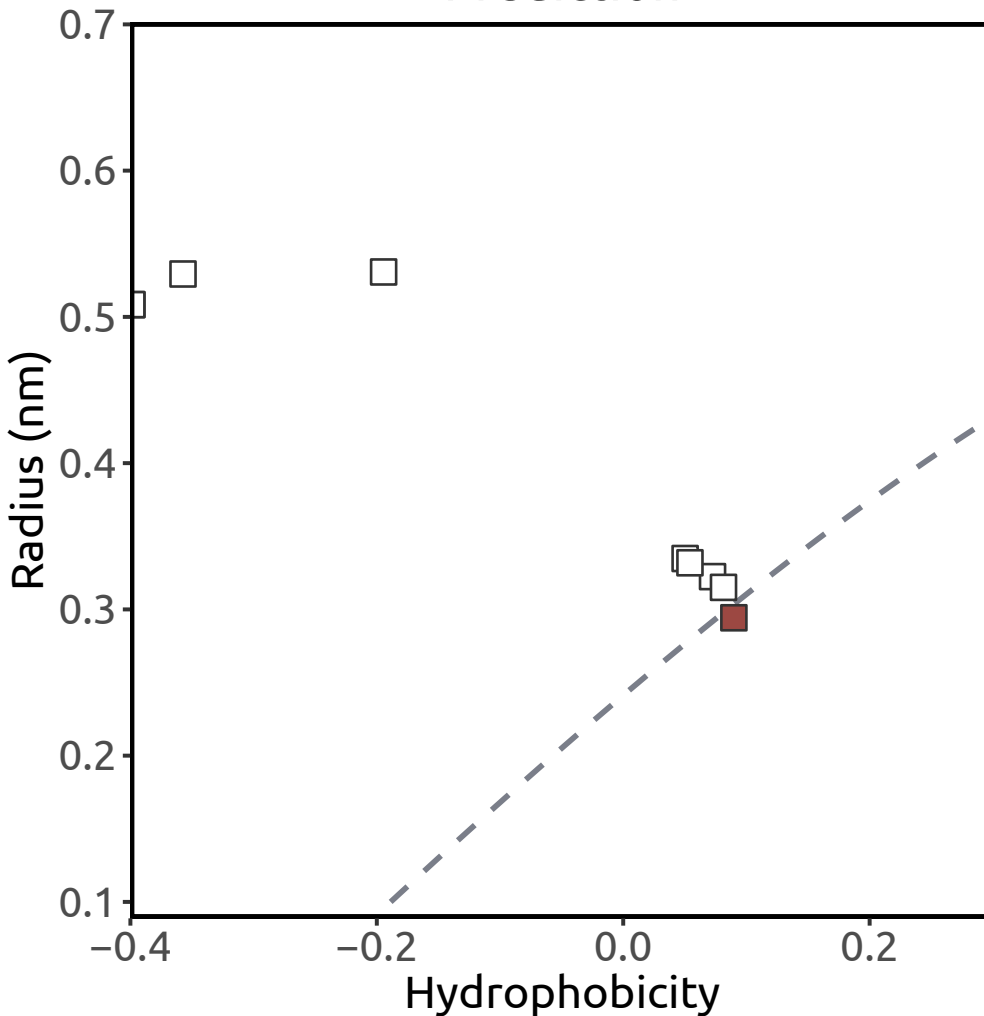
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.01 (n = 1)

Simulation result:
hydrated channel

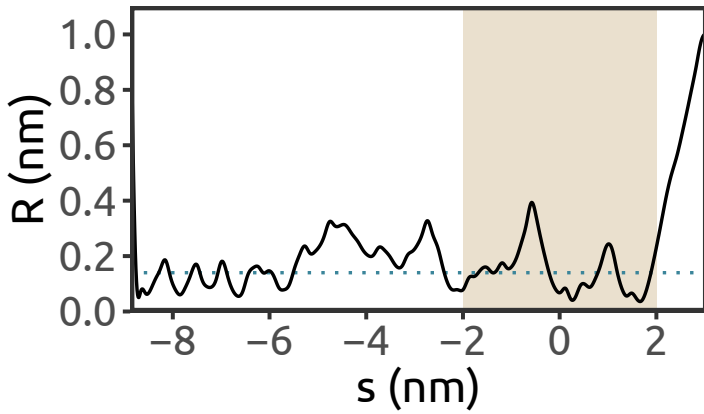
NaKNav (PDB ID: 3VOU)

B. mycooides S. pontiacus

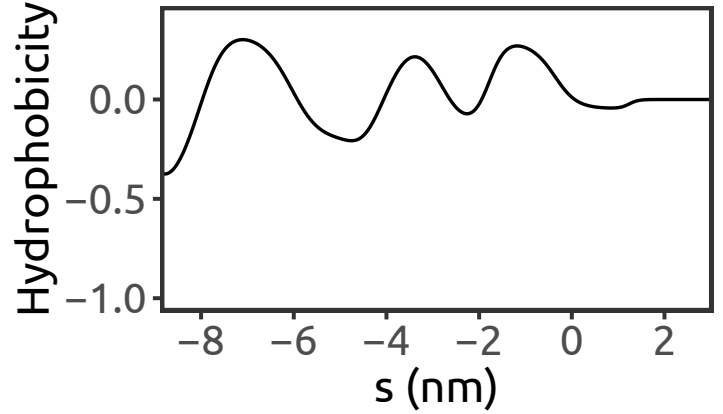
X-ray (3.2 Å)

Irie et al., 2012

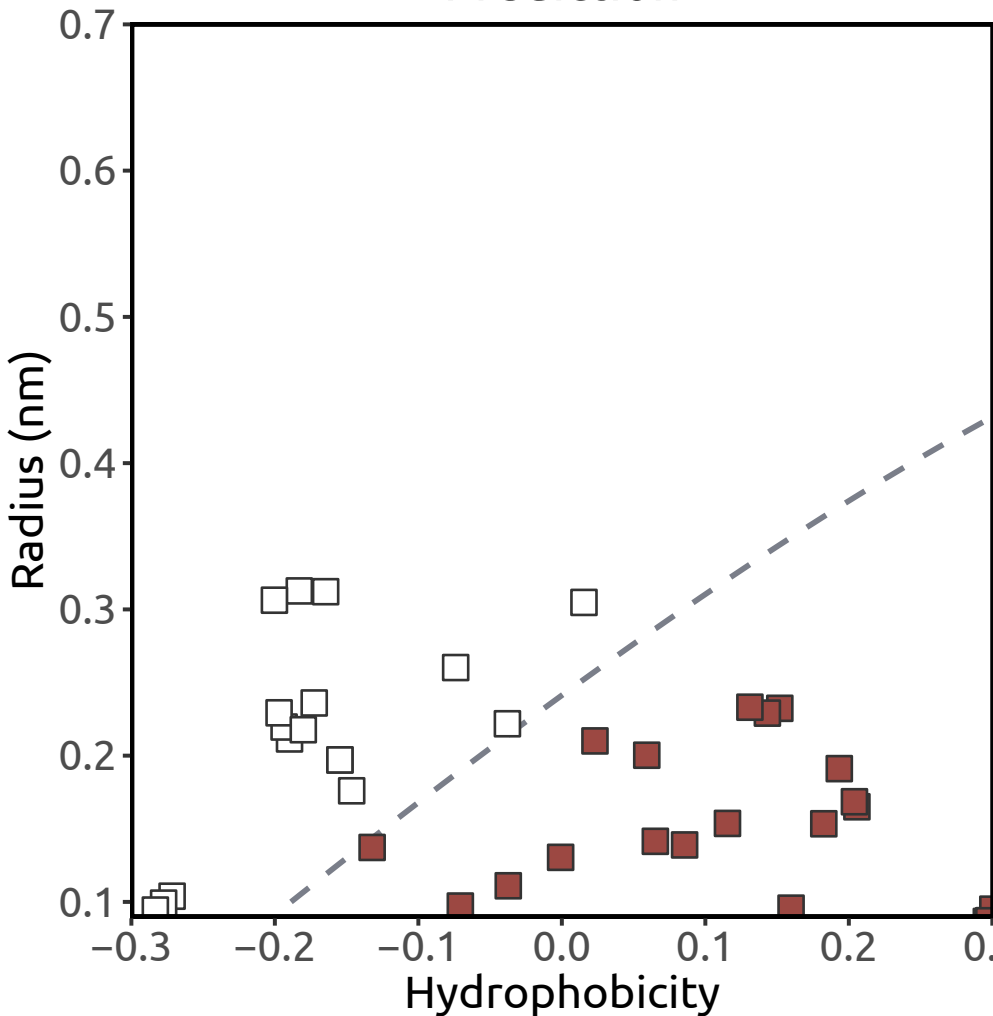
Pore radius



Hydrophobicity



Prediction



Heuristic score:
4.81 (n = 31)

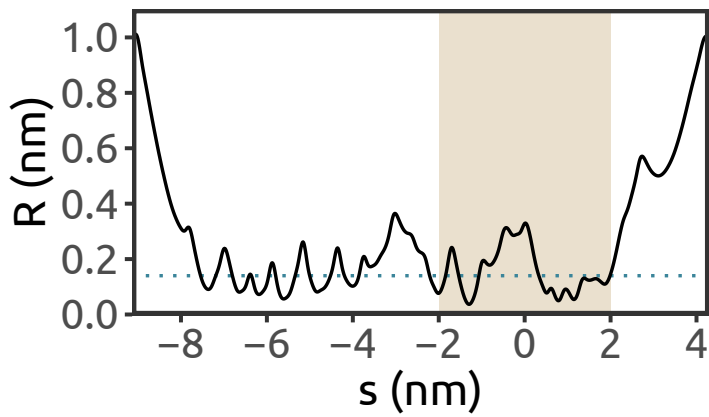
Simulation result:
barrier to water

KCNQ1 (PDB ID: 5VMS)

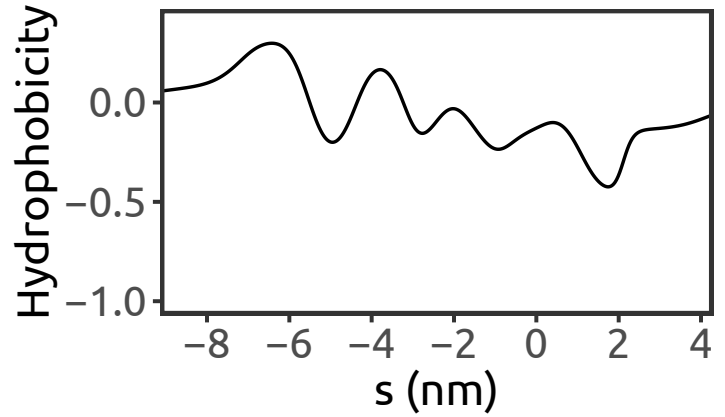
Xenopus laevis
cryo-EM (3.7 Å)

Sun & MacKinnon, 2017

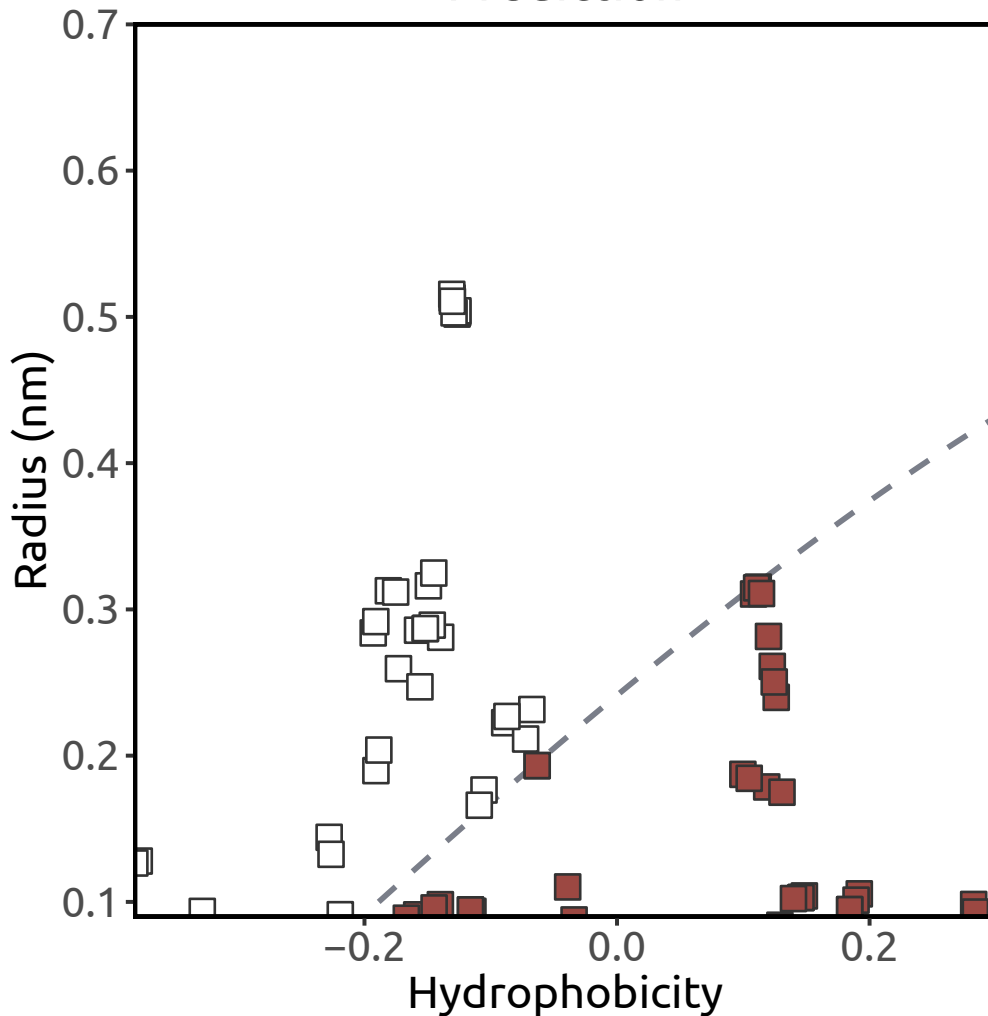
Pore radius



Hydrophobicity



Prediction



Heuristic score:
6.44 (n = 50)

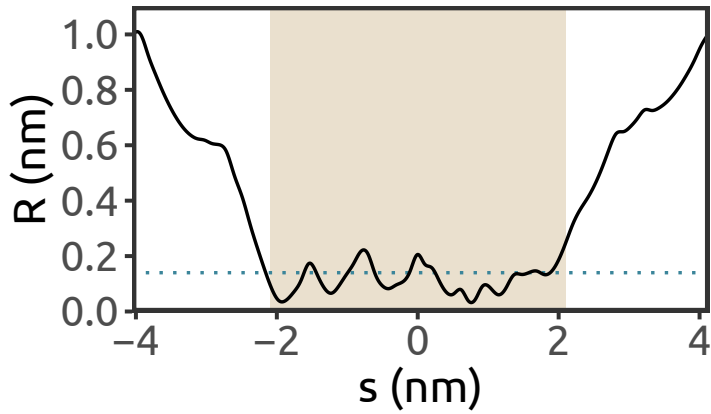
Simulation result:
barrier to water

Kv10.1 (PDB ID: 5K7L)

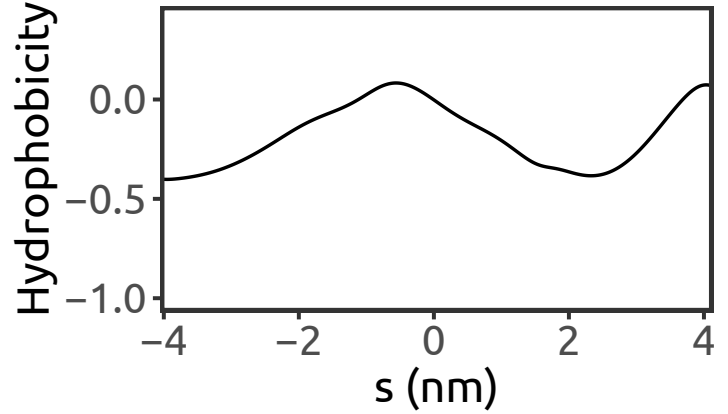
Rattus norvegicus
cryo-EM (3.78 Å)

Whicher & MacKinnon, 2016

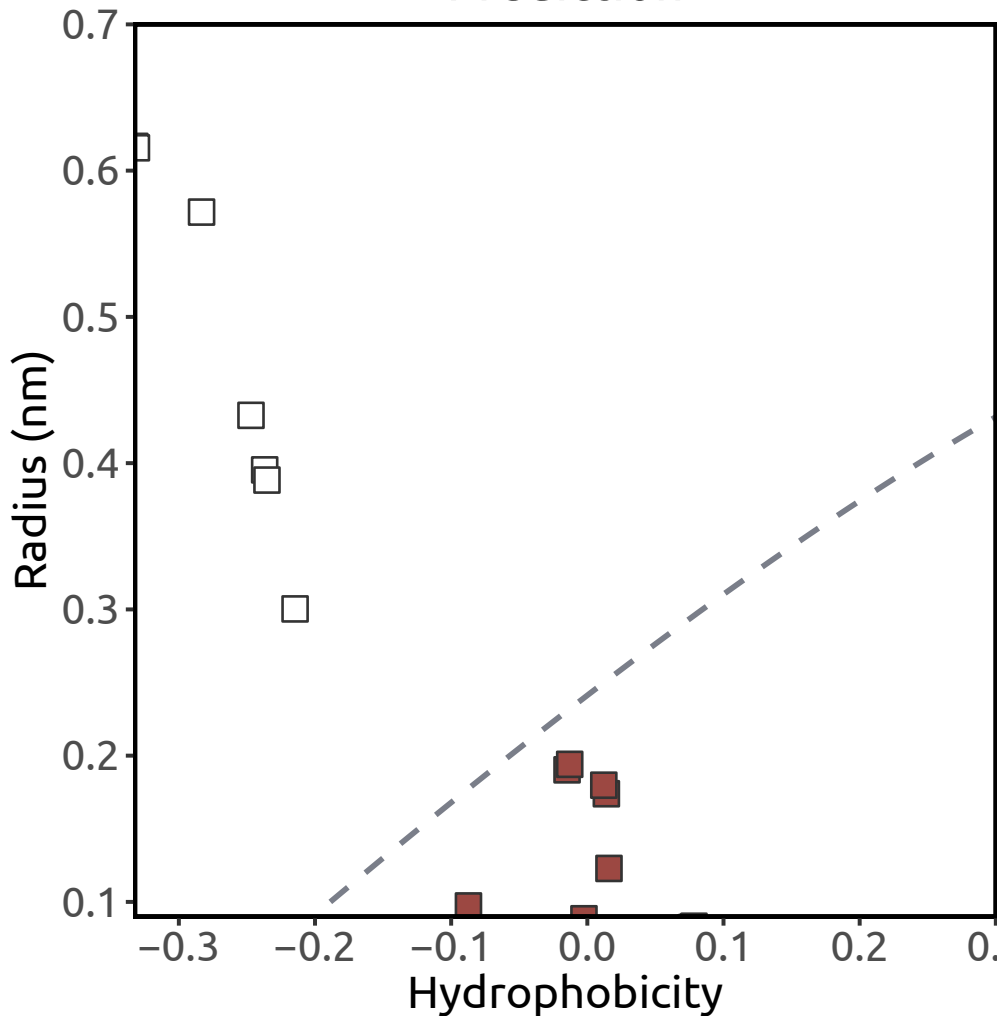
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.69 (n = 16)

Simulation result:
barrier to water

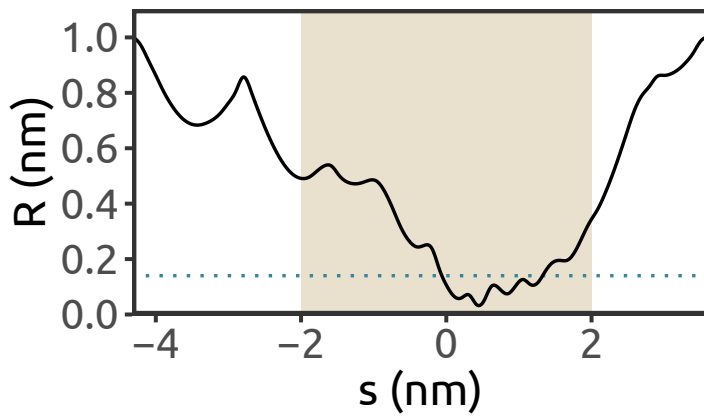
Kv11.1 (PDB ID: 5VA1)

Homo sapiens

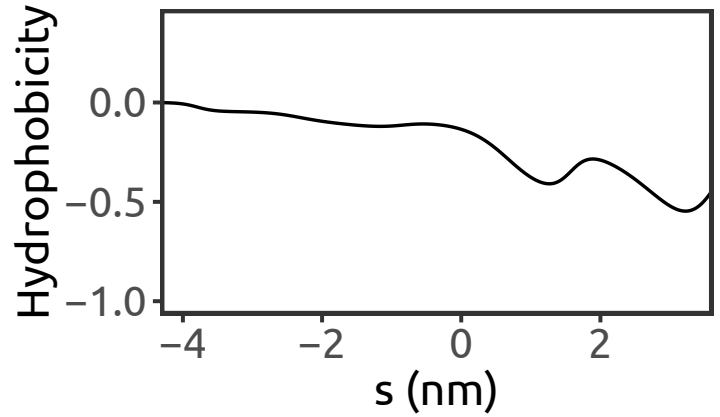
cryo-EM (3.7 Å)

Wang & MacKinnon, 2017

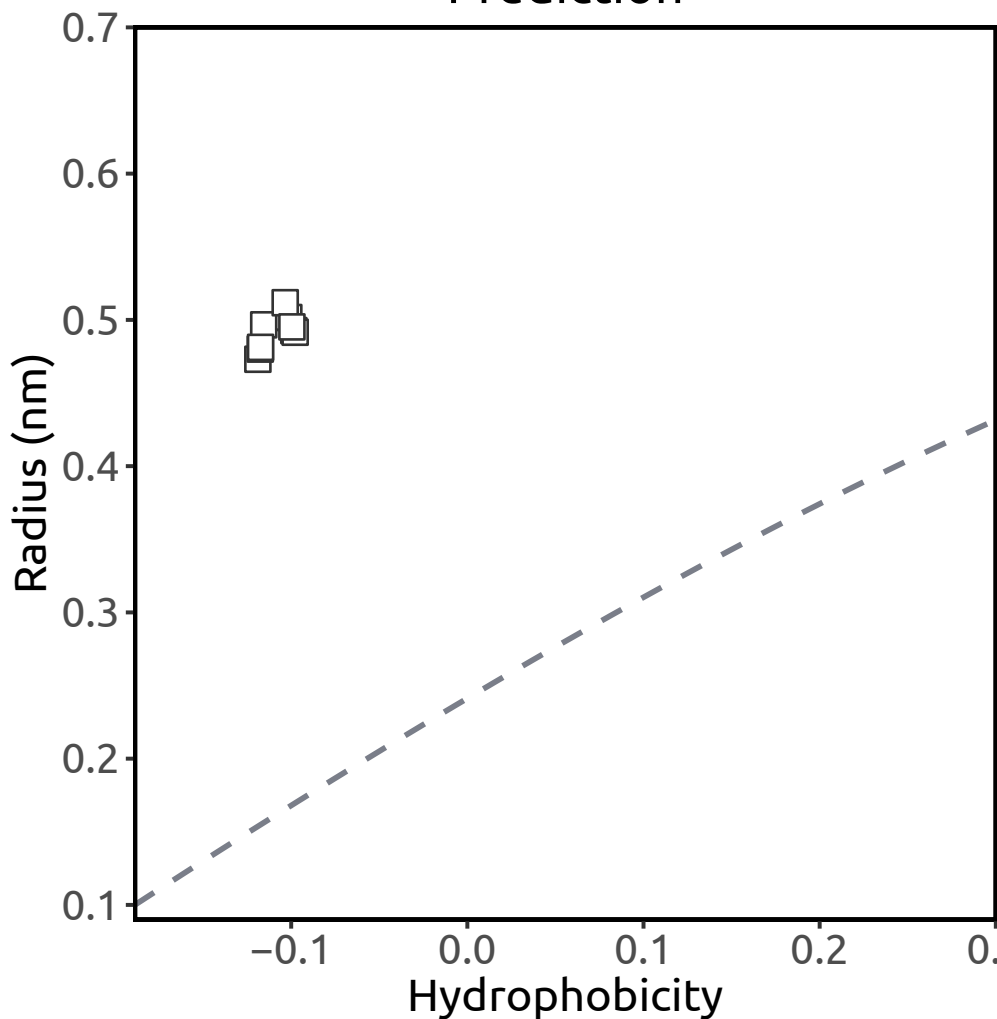
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

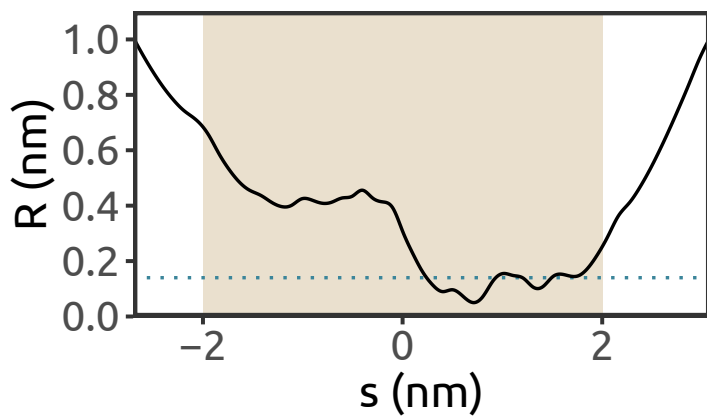
Kv1.2 (PDB ID: 2A79)

Rattus norvegicus

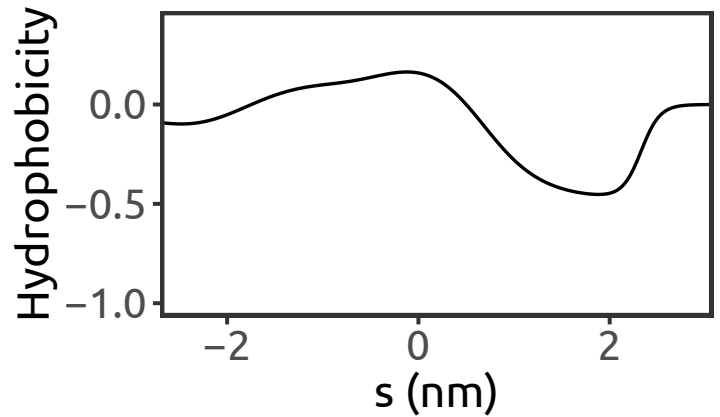
X-ray (2.9 Å)

Long et al., 2005

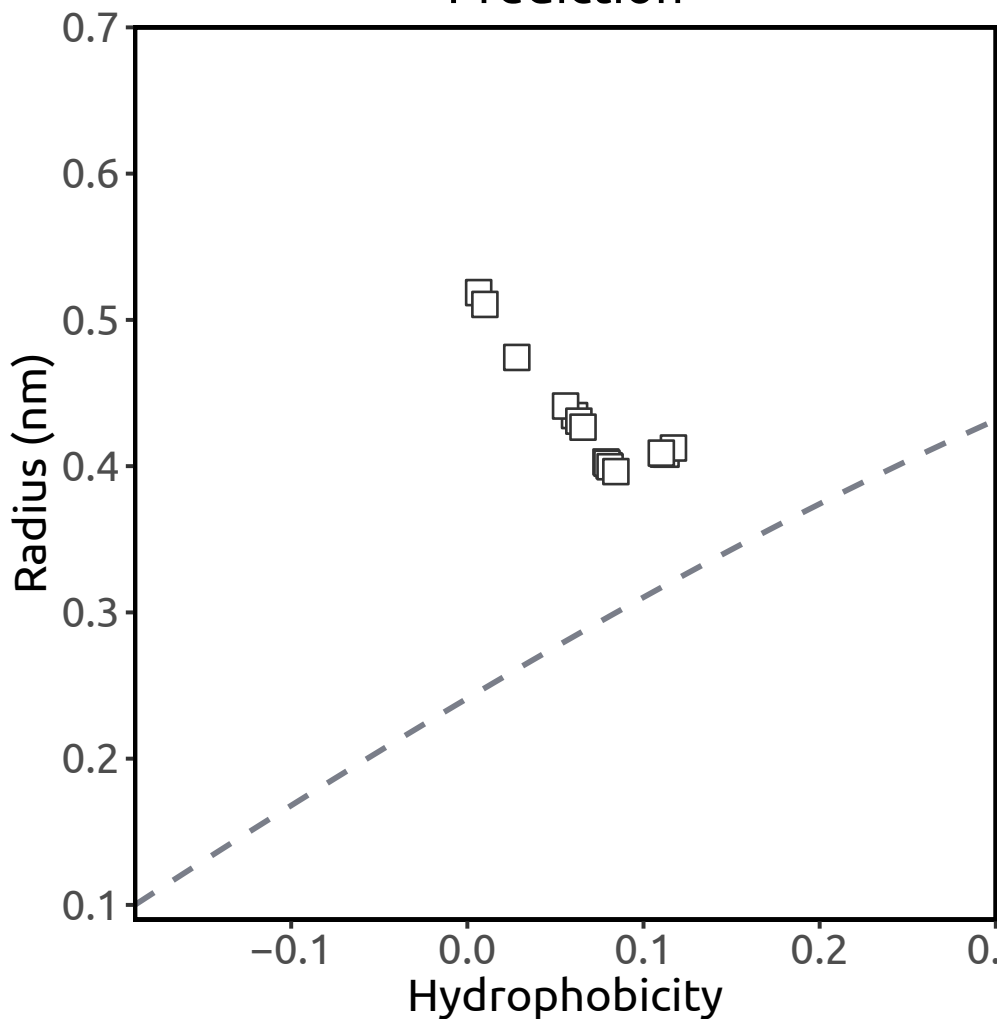
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

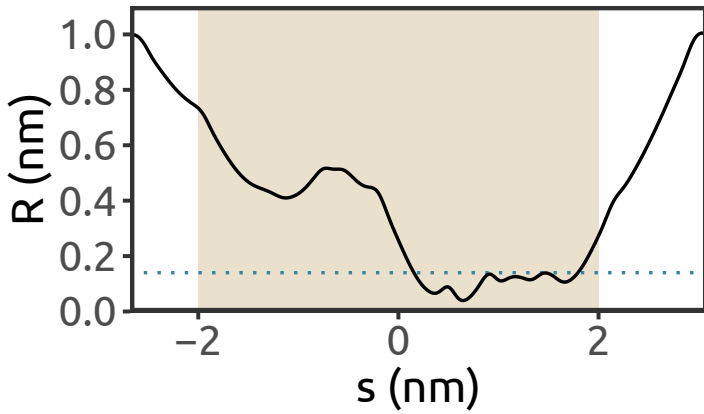
Kv1.2 (PDB ID: 2R9R)

Rattus norvegicus

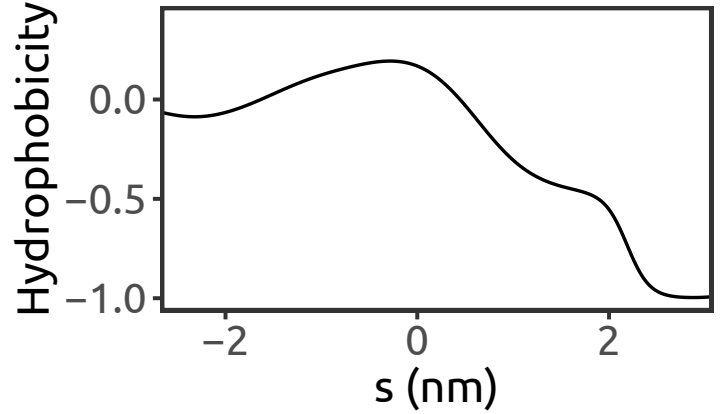
X-ray (2.4 Å)

Long et al., 2007

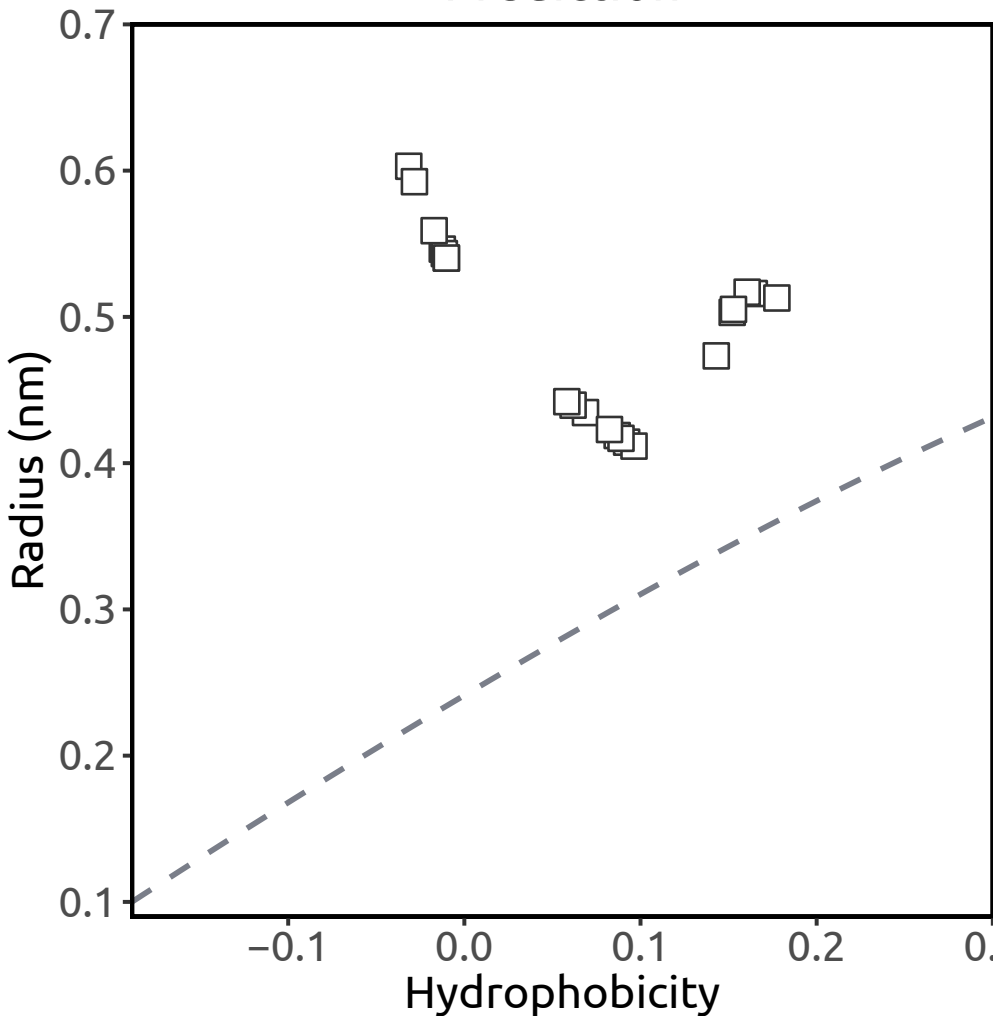
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 ($n = 0$)

Simulation result:
hydrated channel

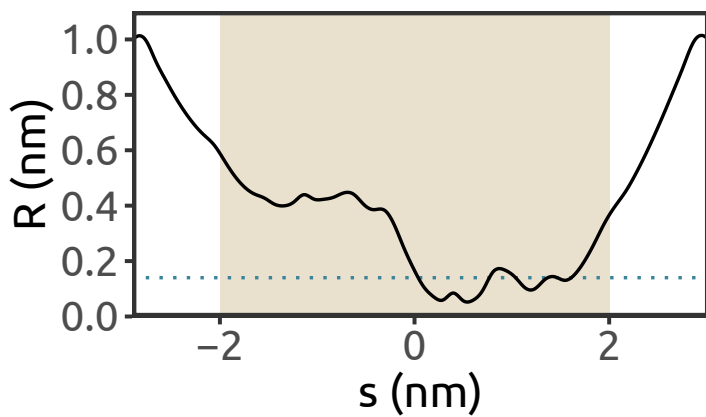
Kv1.2 (PDB ID: 3LUT)

Rattus norvegicus

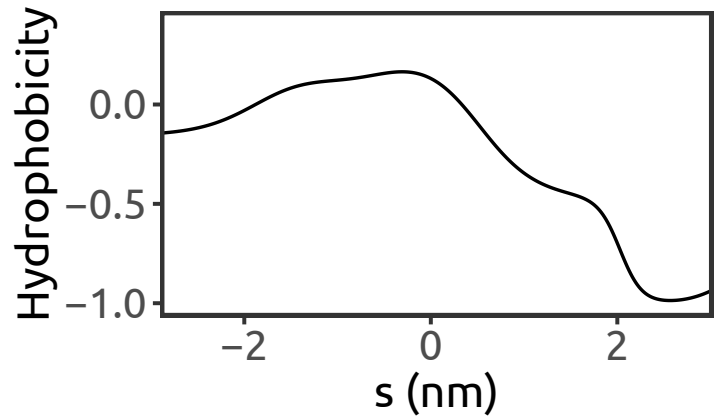
X-ray (2.9 Å)

Chen et al., 2010

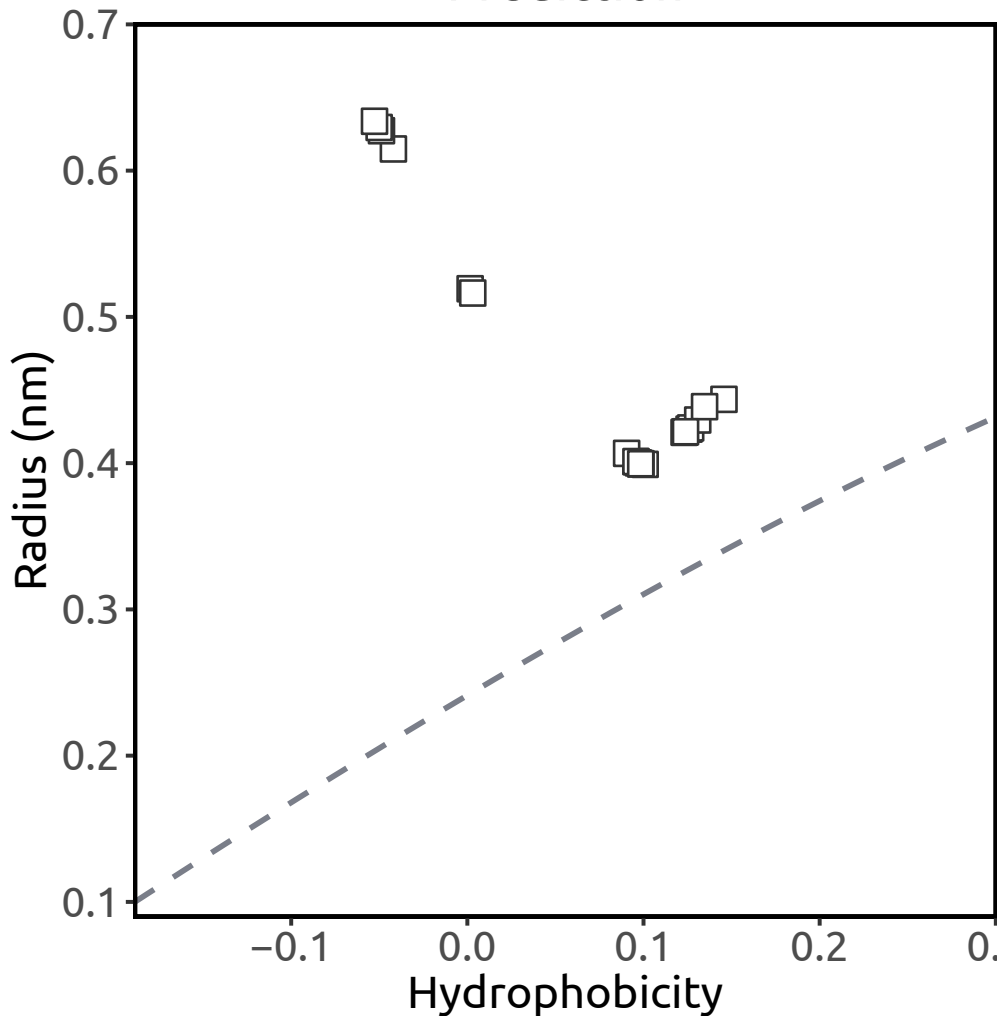
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

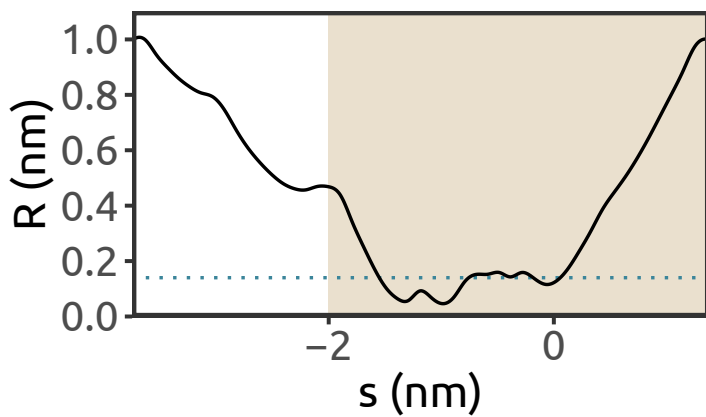
KvAp (PDB ID: 1ORQ)

Aeropyrum pernix

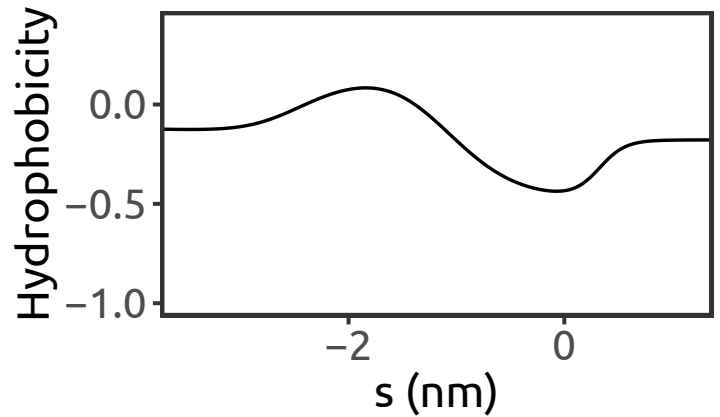
X-ray (3.2 Å)

Jiang et al., 2003

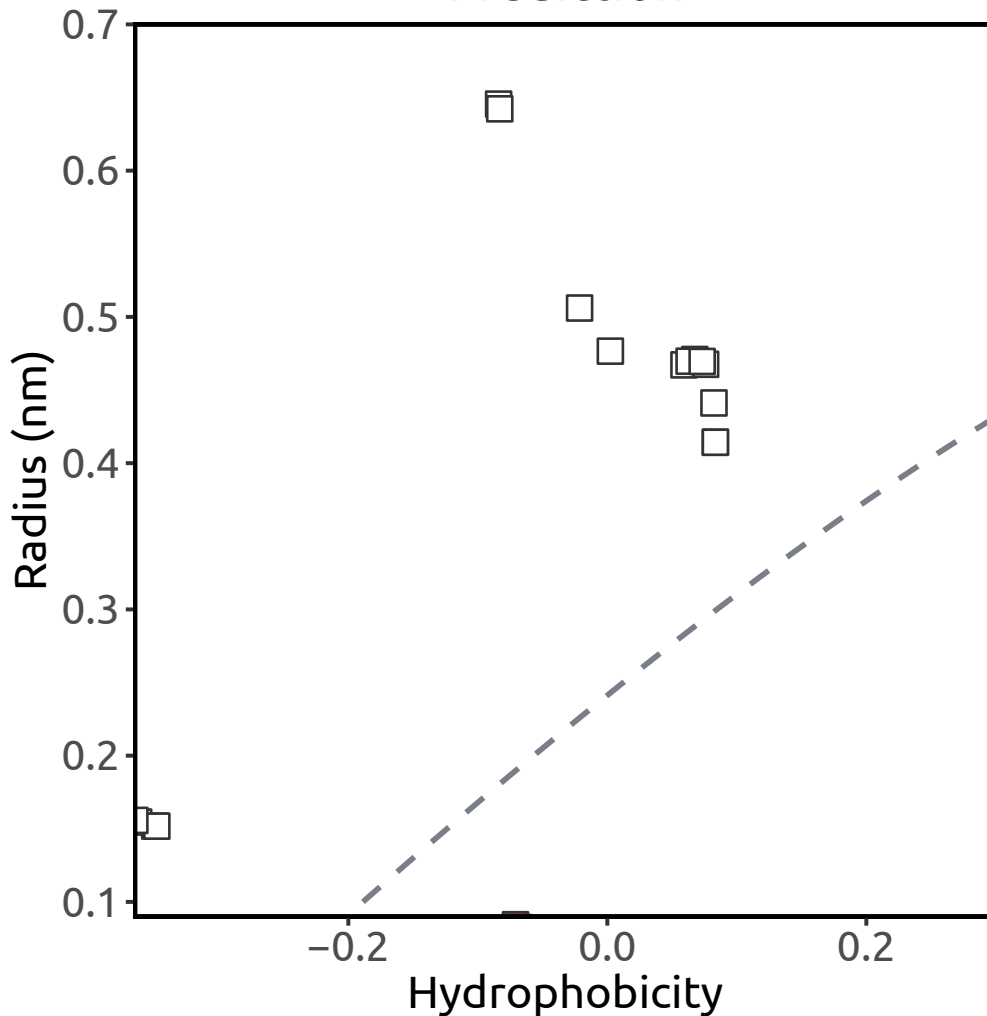
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.38 (n = 4)

Simulation result:
barrier to water

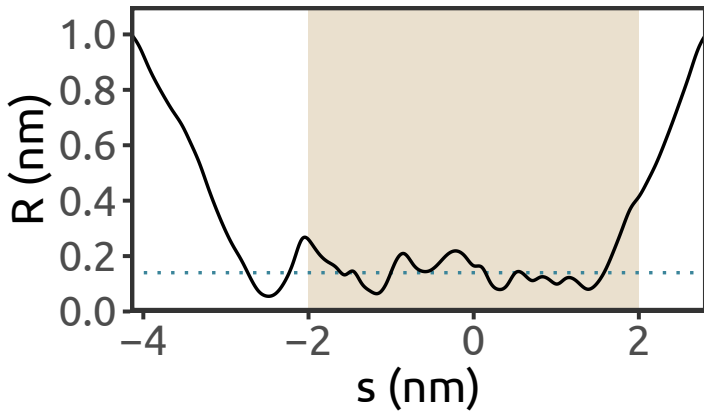
KvLm (PDB ID: 4H33)

Listeria monocytogenes

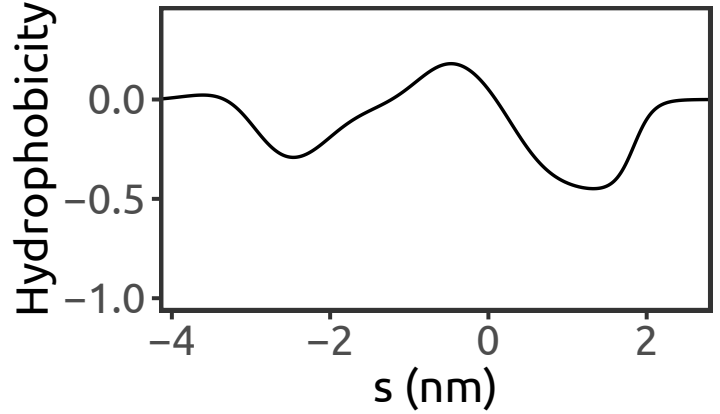
X-ray (3.1 Å)

Santos et al., 2012

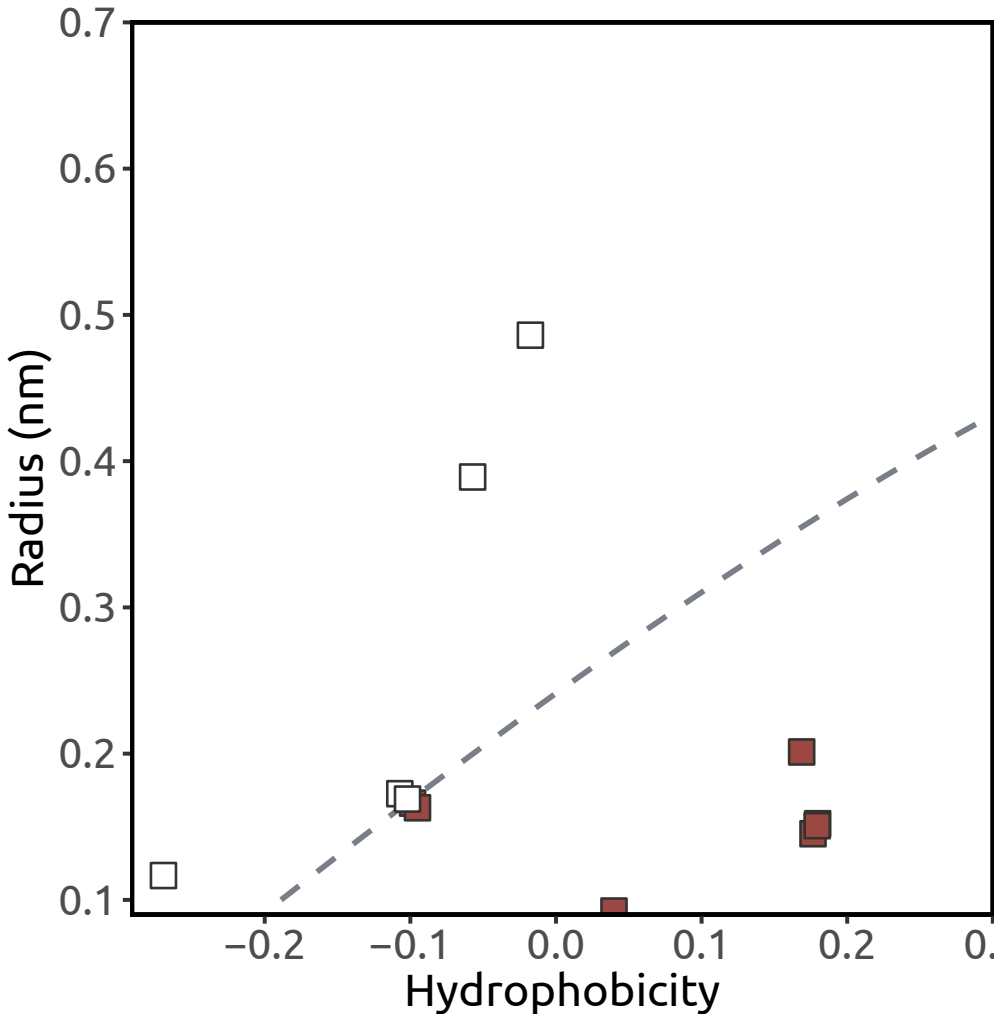
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.25 (n = 10)

Simulation result:
barrier to water

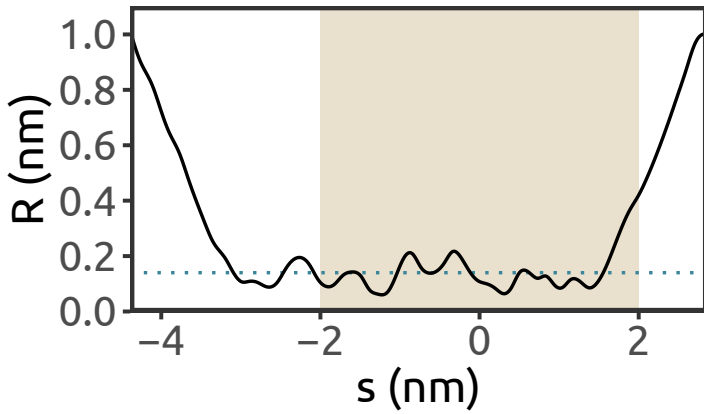
KvLm (PDB ID: 4H37)

Listeria monocytogenes

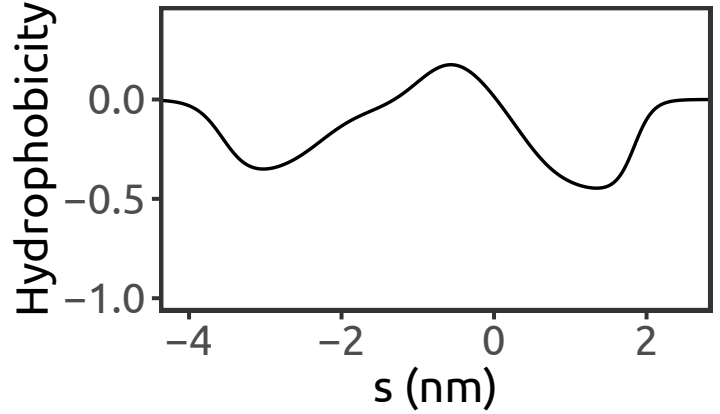
X-ray (3.35 Å)

Santos et al., 2012

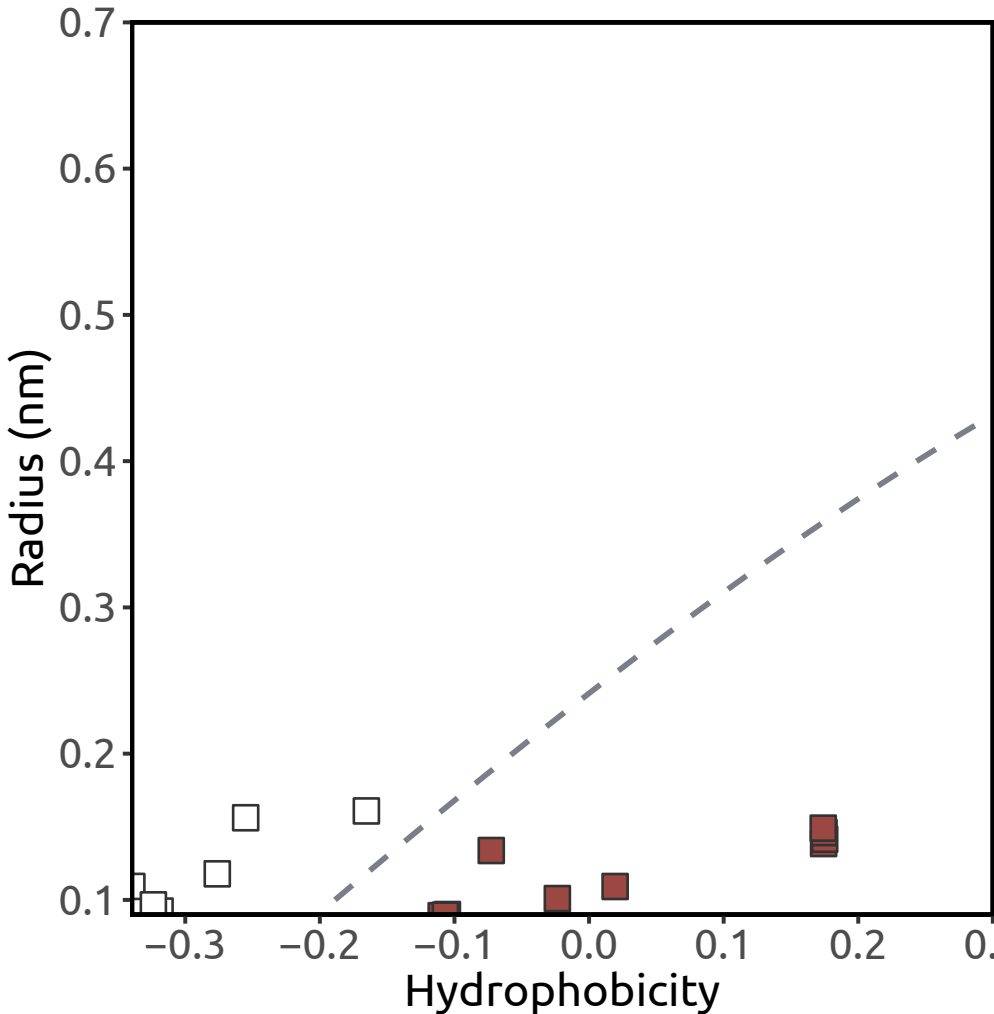
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.76 (n = 14)

Simulation result:

barrier to water

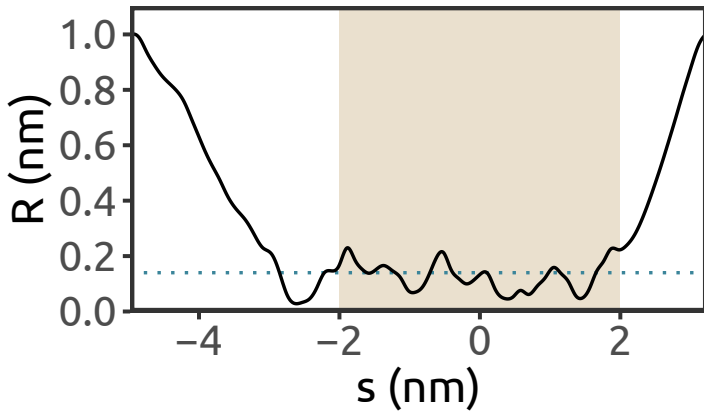
MloK1 (PDB ID: 3BEH)

Mesorhizobium loti

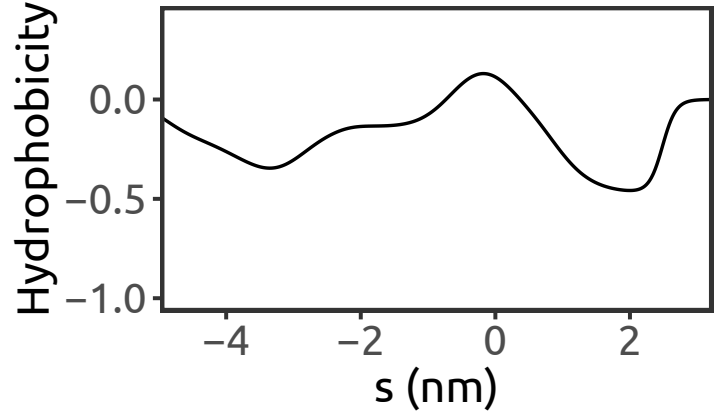
X-ray (3.1 Å)

Clayton et al., 2008

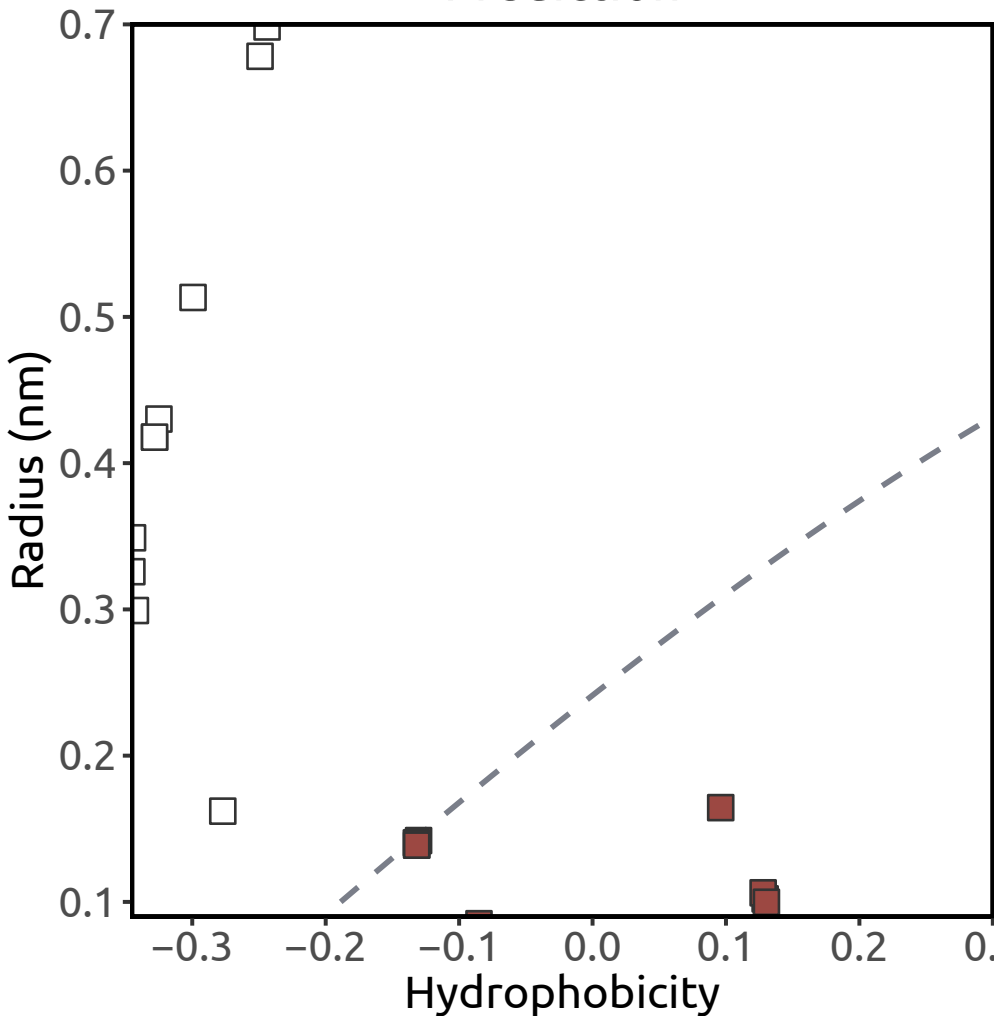
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.13 (n = 13)

Simulation result:

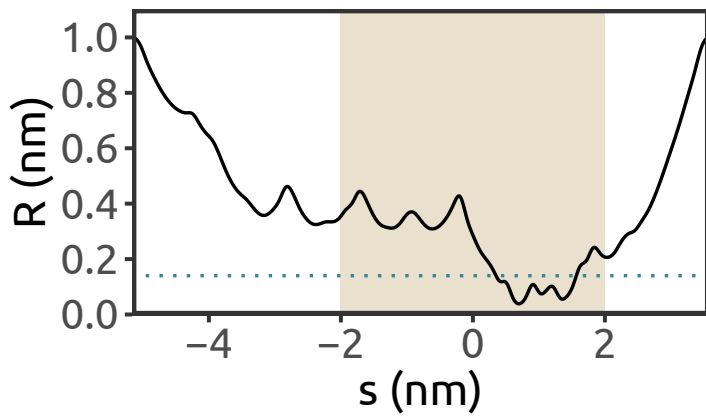
barrier to water

MloK1 (PDB ID: 6EO1)

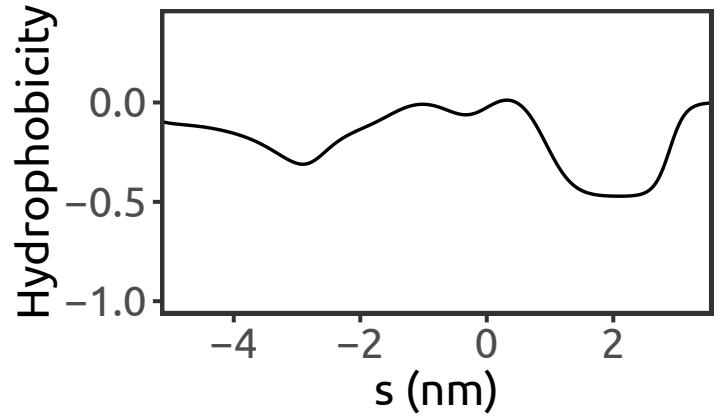
Mesorhizobium loti
cryo-EM (4.5 Å)

Kowal et al., 2018

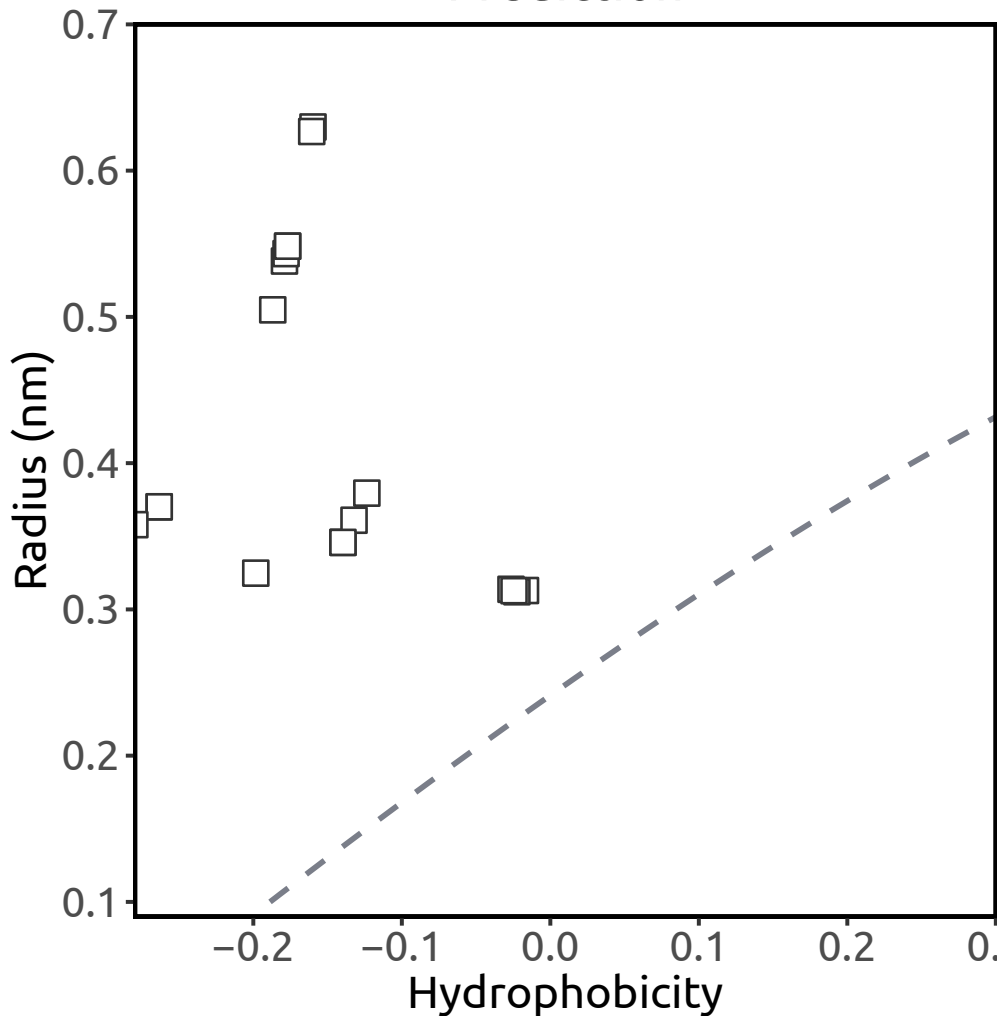
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

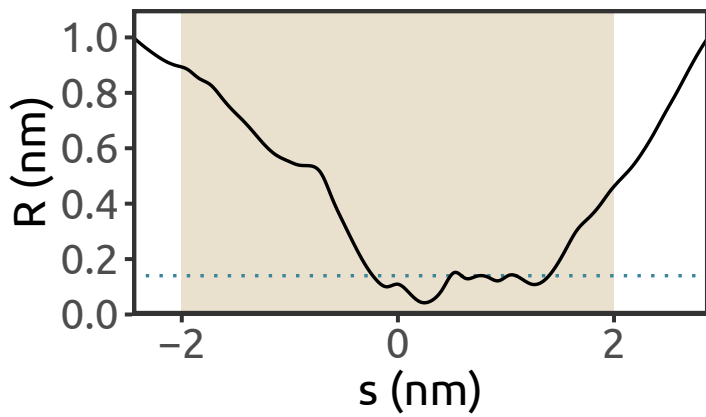
Simulation result:
hydrated channel

Slo1 (PDB ID: 5TJ6)

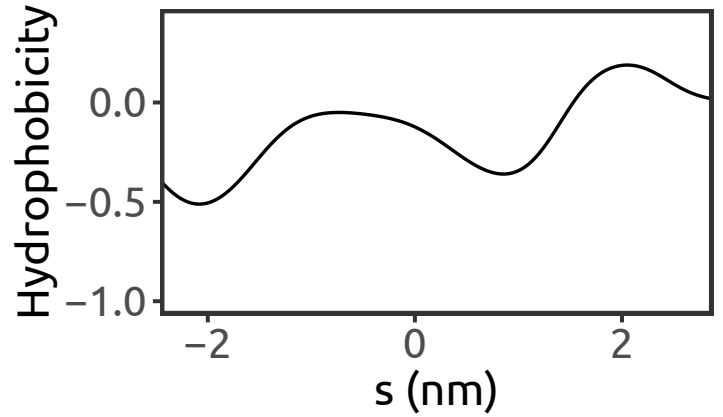
Aplysia californica
cryo-EM (3.5 Å)

Tao et al, 2017

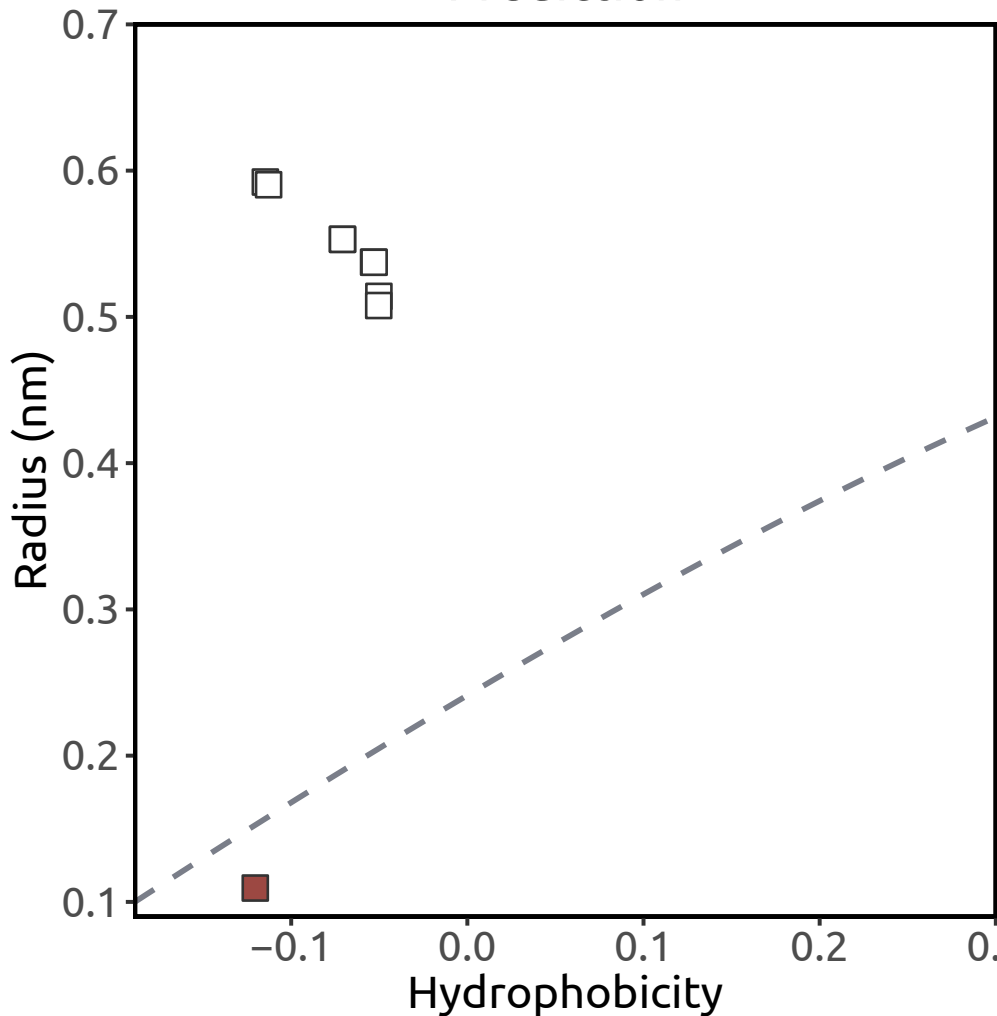
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.03 (n = 1)

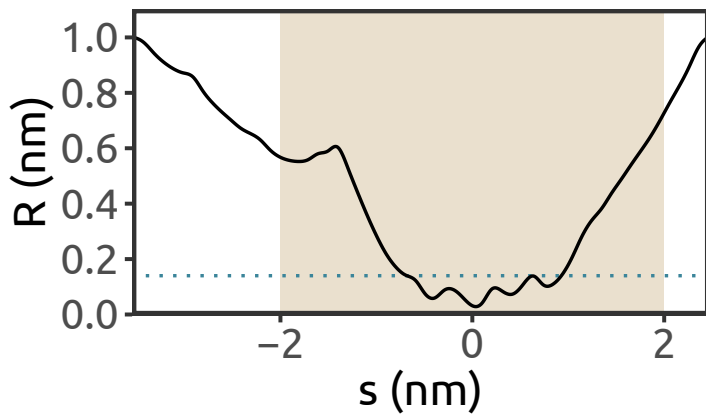
Simulation result:
hydrated channel

Slo1 (PDB ID: 5TJI)

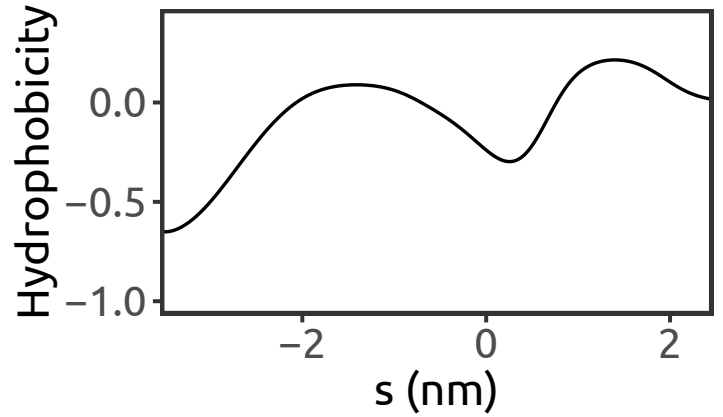
Aplysia californica
cryo-EM (3.8 Å)

Hite et al., 2017

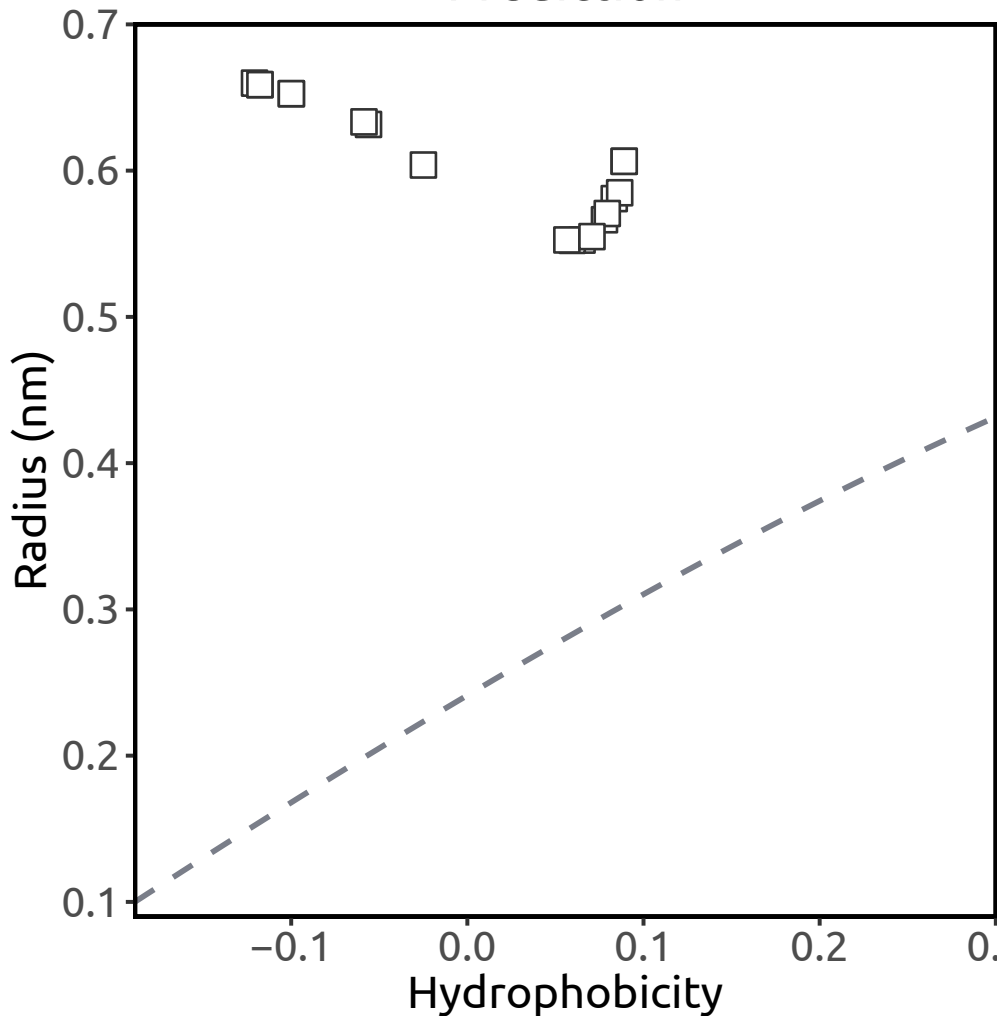
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

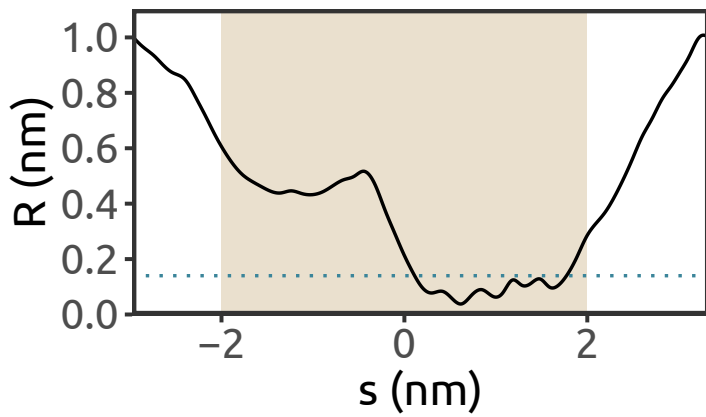
Slo2.2 (PDB ID: 5U70)

Gallus gallus

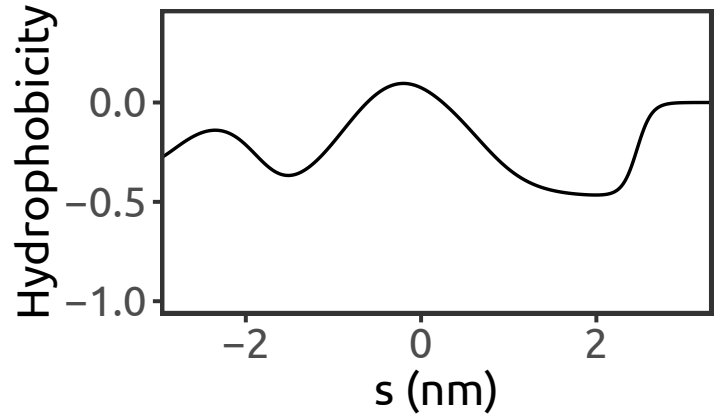
cryo-EM (3.76 Å)

Hite & MacKinnon, 2017

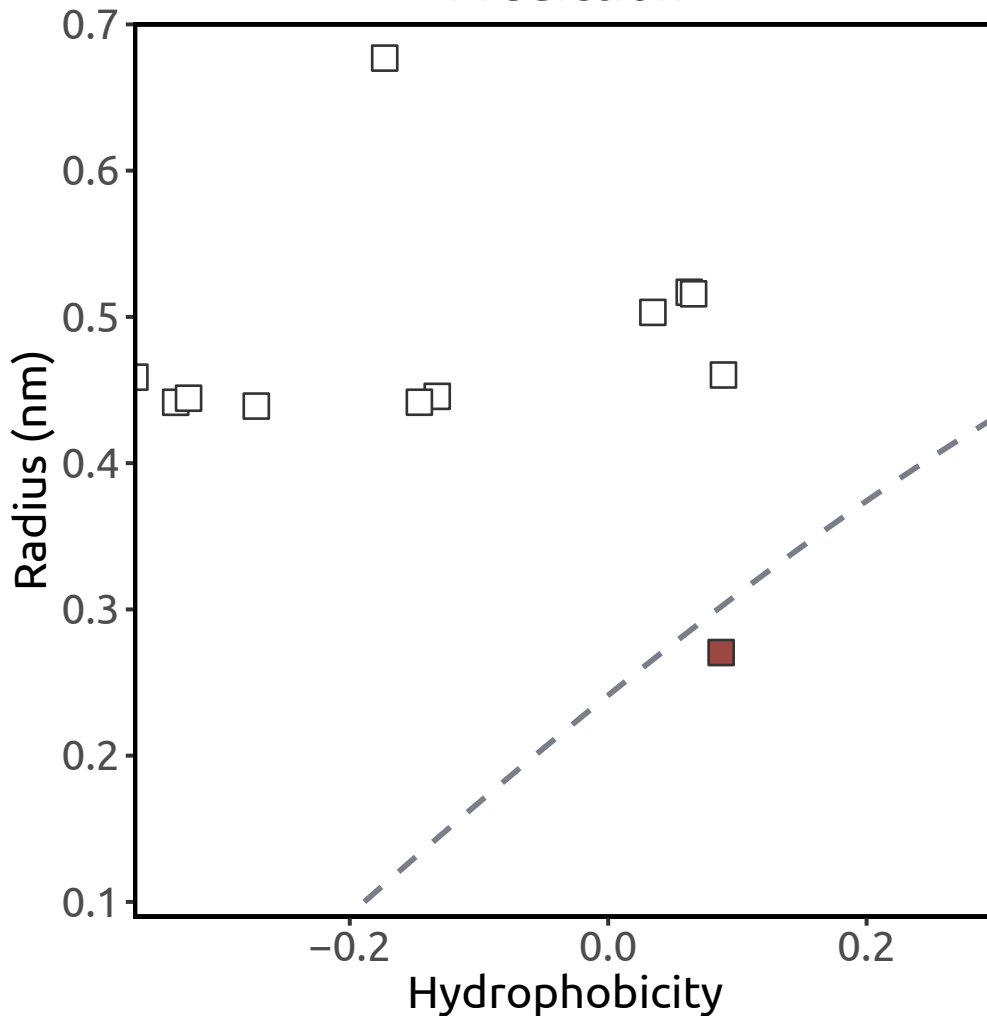
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.03 ($n = 1$)

Simulation result:
hydrated channel

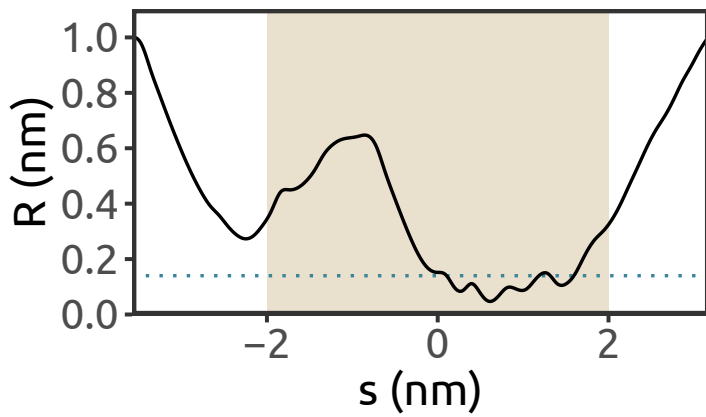
Slo2.2 (PDB ID: 5U76)

Gallus gallus

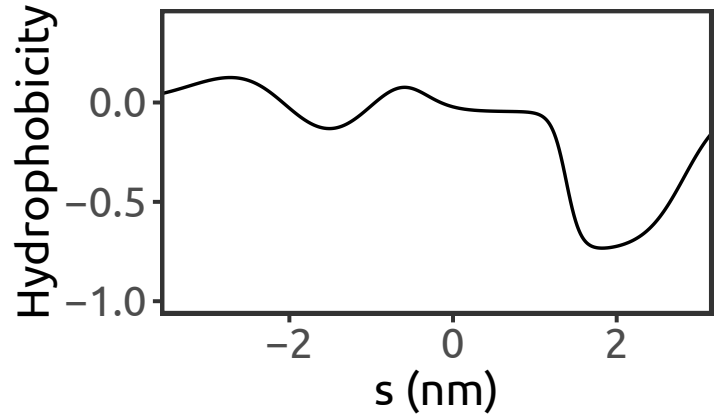
cryo-EM (3.76 Å)

Hite & MacKinnon, 2017

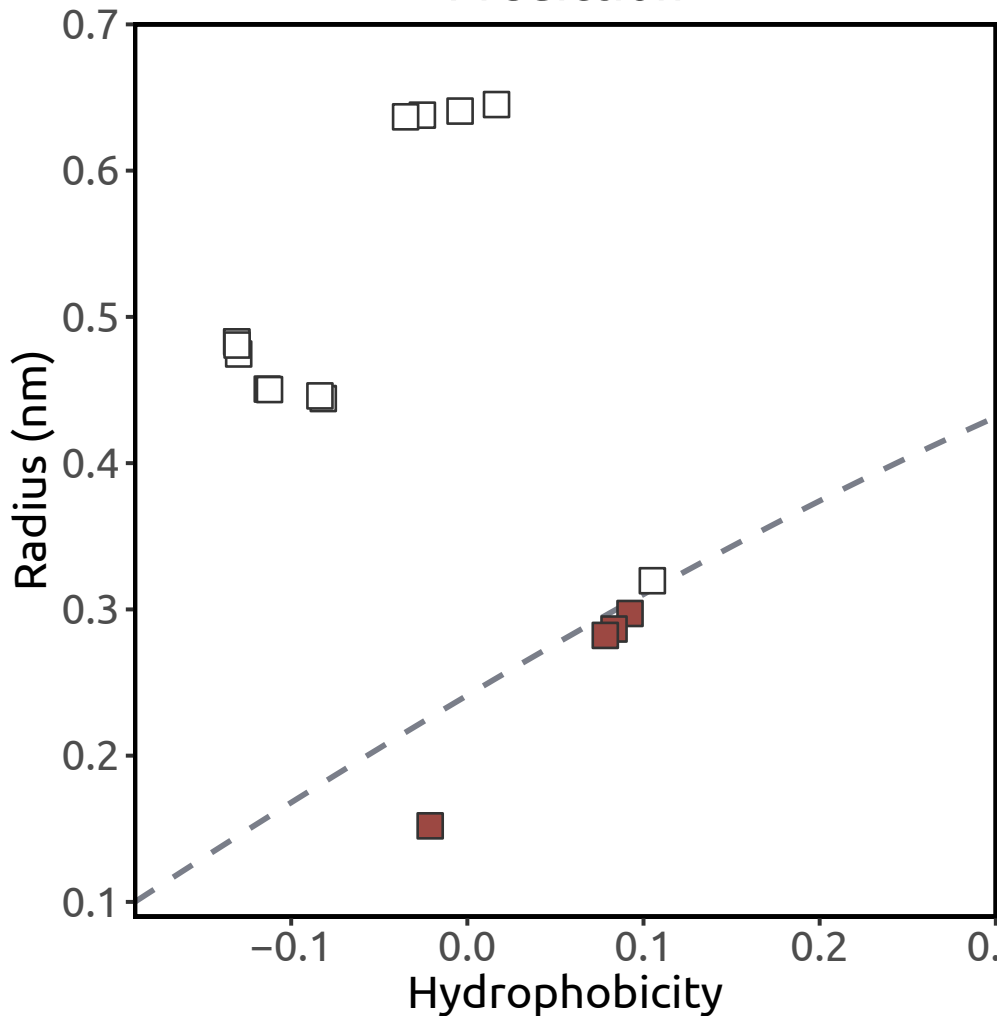
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.09 (n = 4)

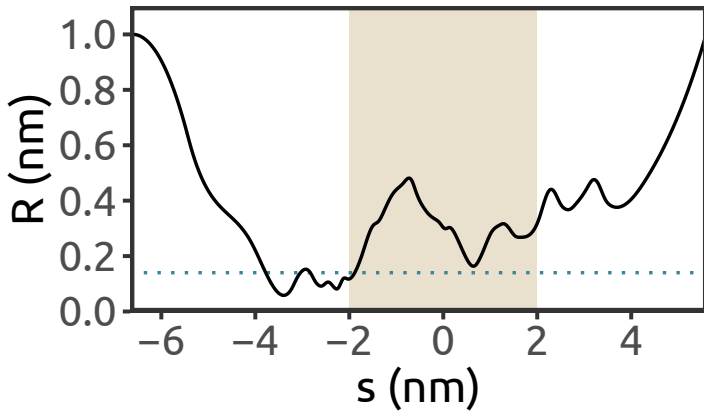
Simulation result:
hydrated channel

Cav1.1 (PDB ID: 5GJV)

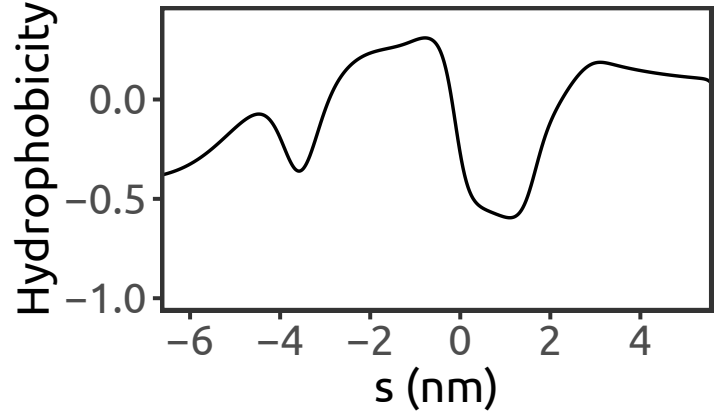
Oryctolagus cuniculus
cryo-EM (3.6 Å)

Wu et al., 2016

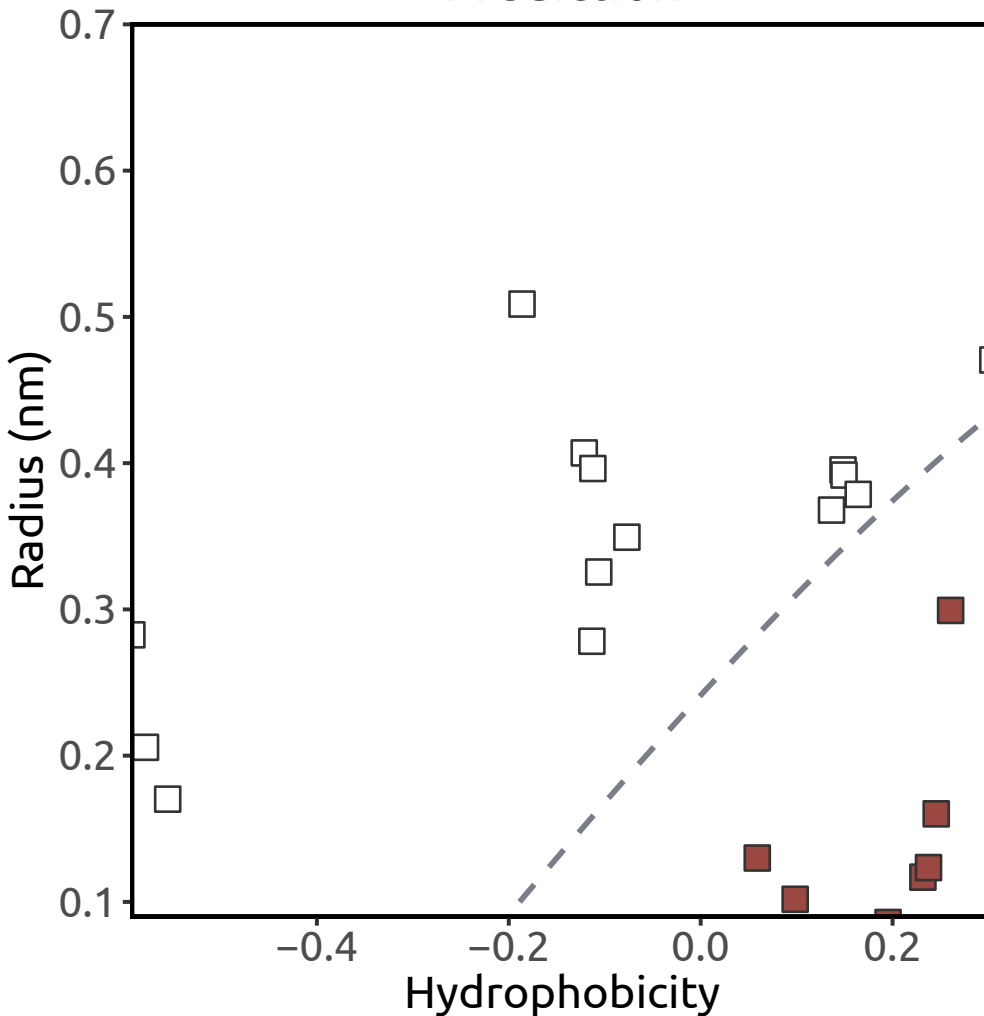
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.3 (n = 7)

Simulation result:
barrier to water

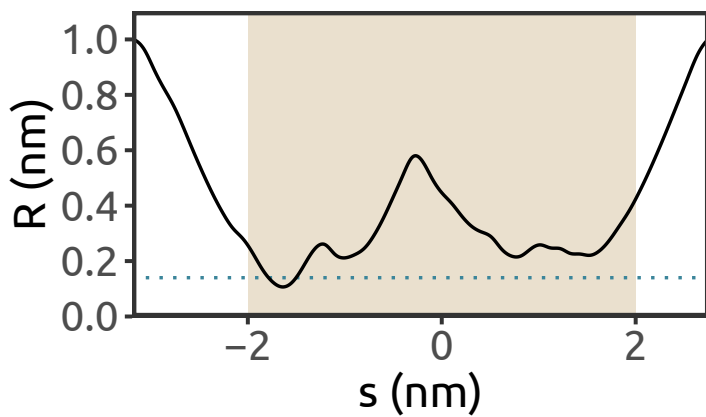
CavAb (PDB ID: 4MVM)

Arcobacter butzleri

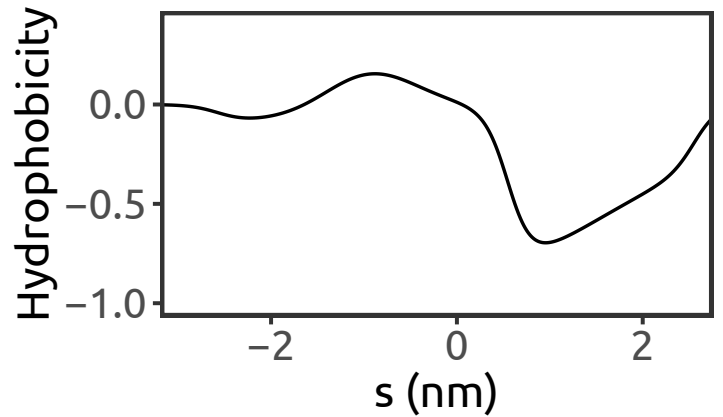
X-ray (3.20 Å)

Tang et al., 2014

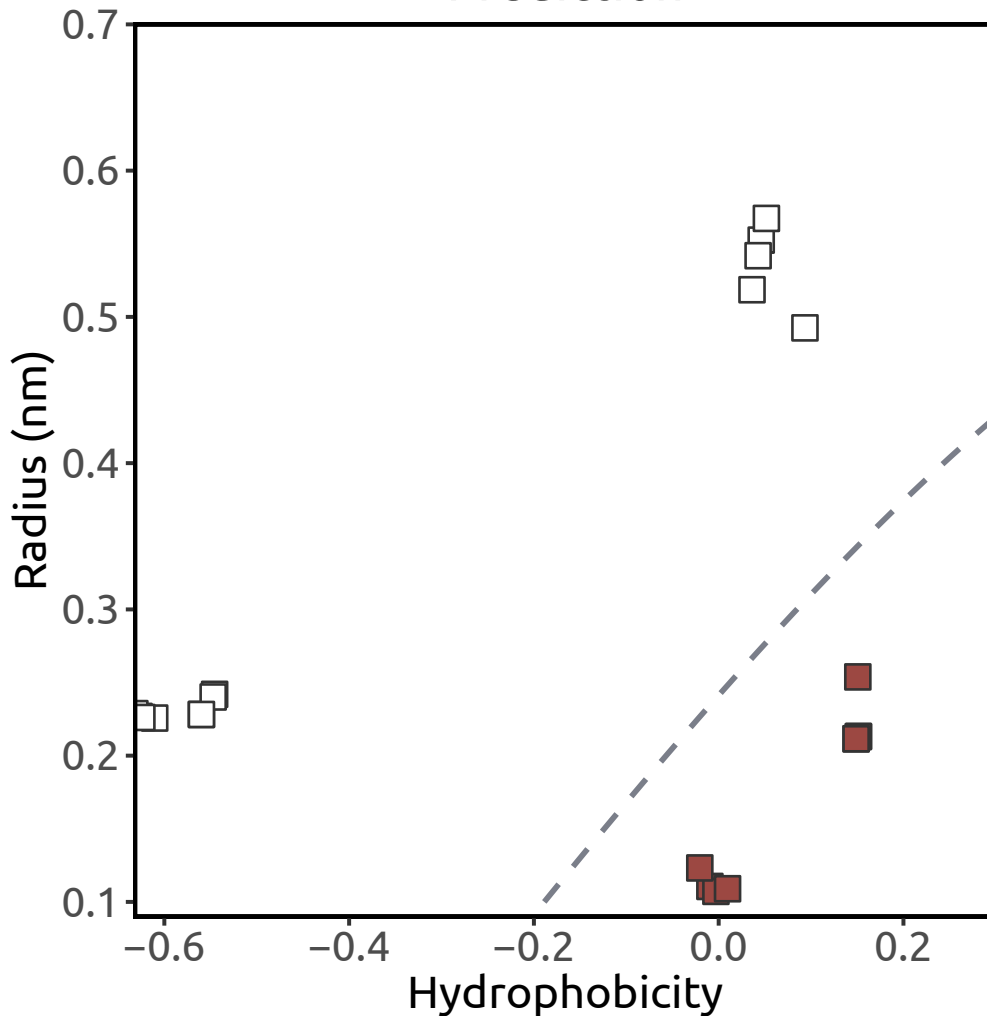
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.92 (n = 9)

Simulation result:
barrier to water

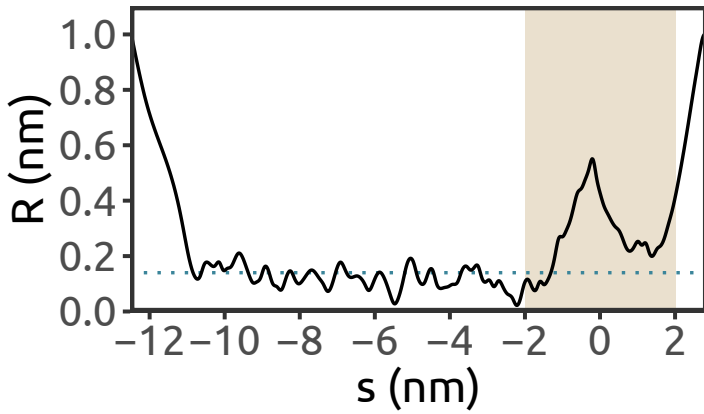
CavAb (PDB ID: 5KLB)

Arcobacter butzleri

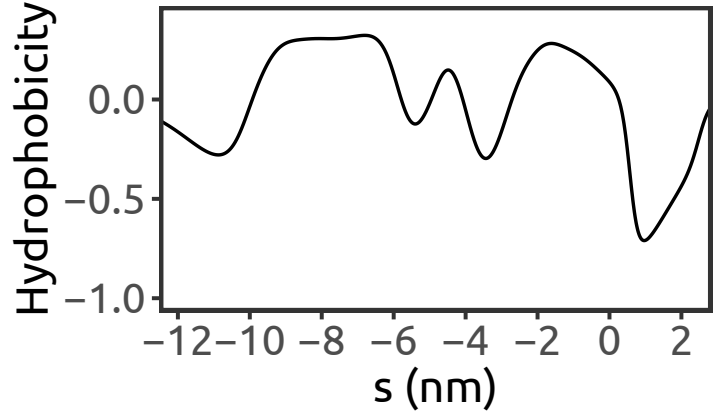
X-ray (2.7 Å)

Tang et al., 2016

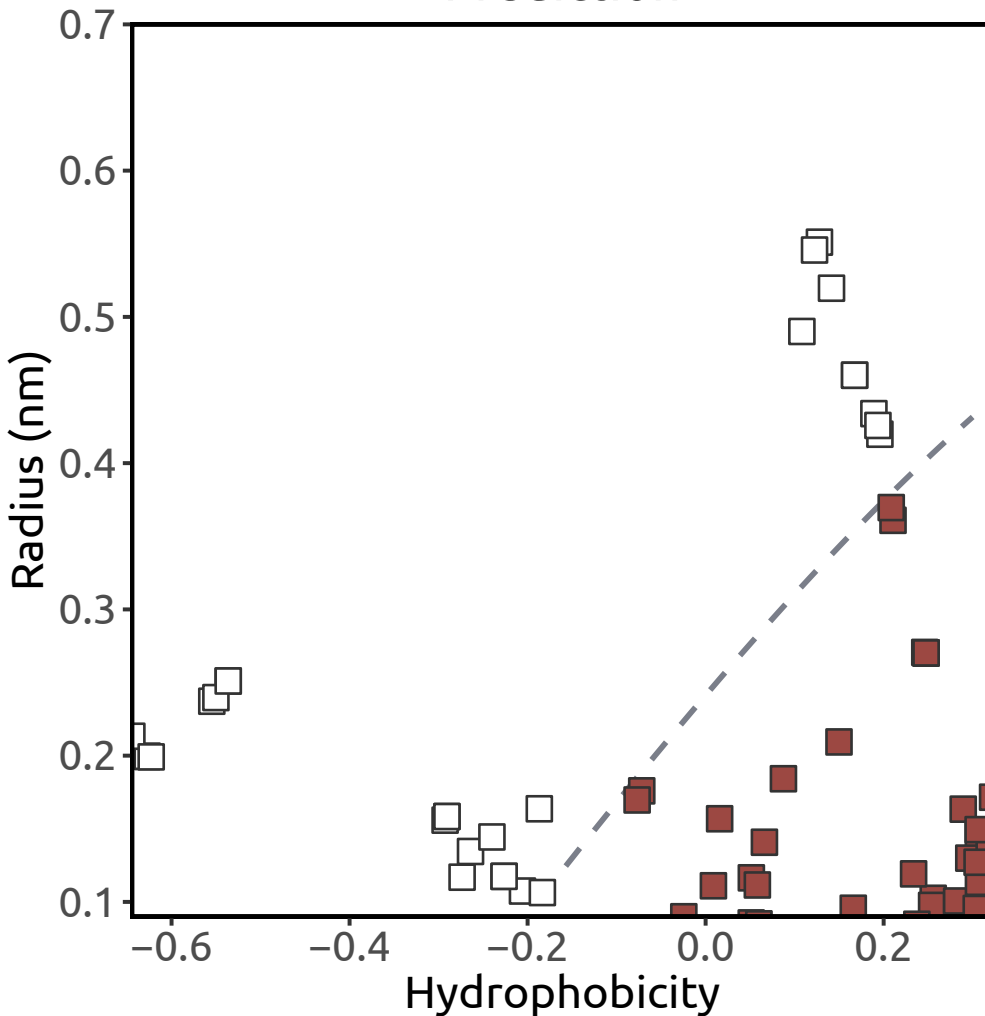
Pore radius



Hydrophobicity



Prediction



Heuristic score:
9.44 (n = 50)

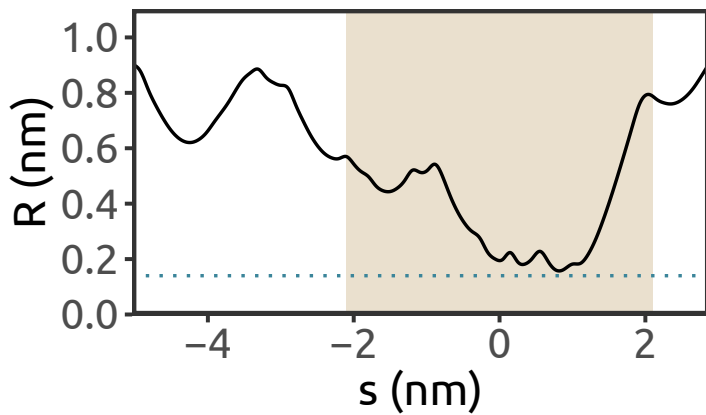
Simulation result:
barrier to water

CNG (PDB ID: 5H3O)

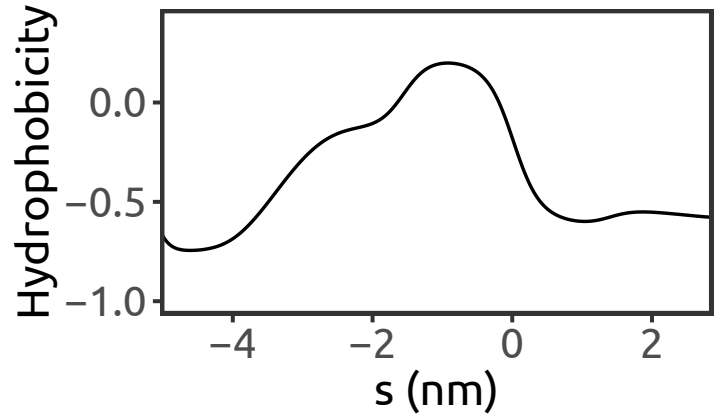
Caenorhabditis elegans
cryo-EM (3.5 Å)

Li et al., 2017

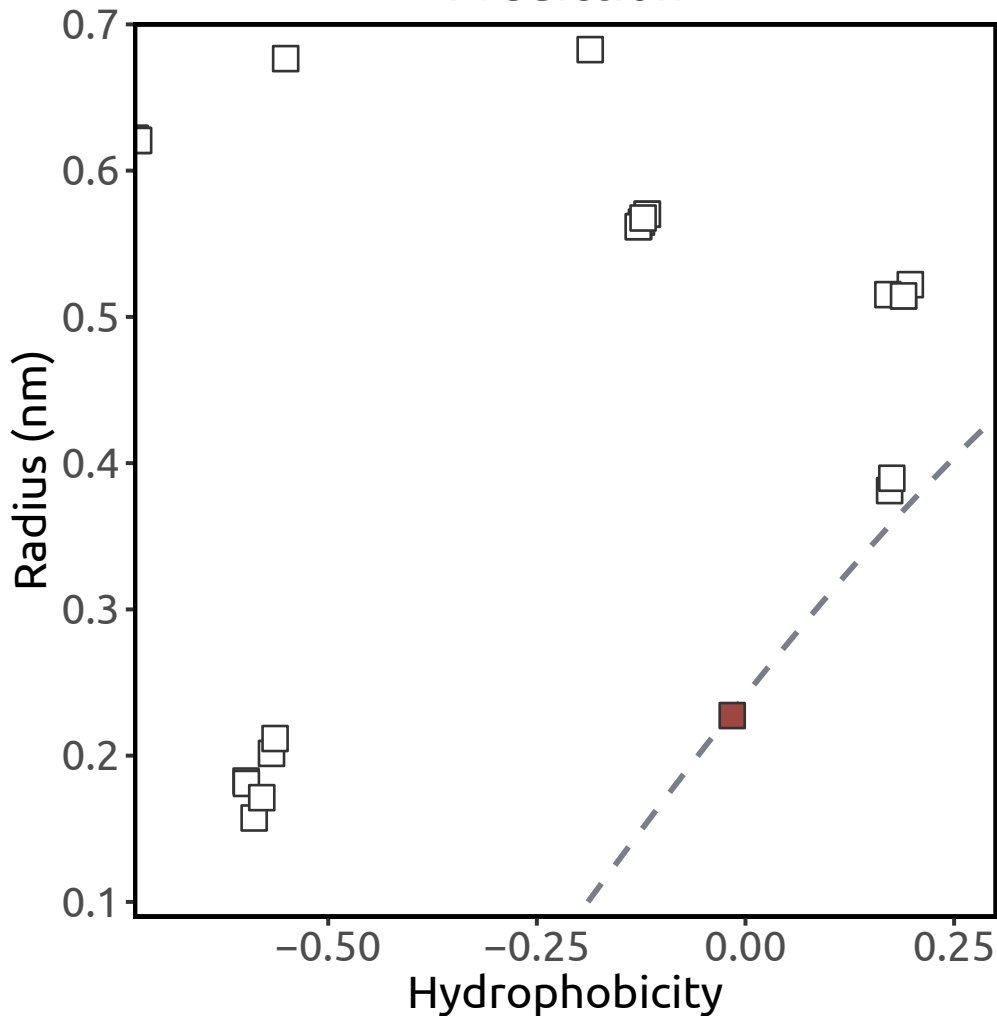
Pore radius



Hydrophobicity



Prediction



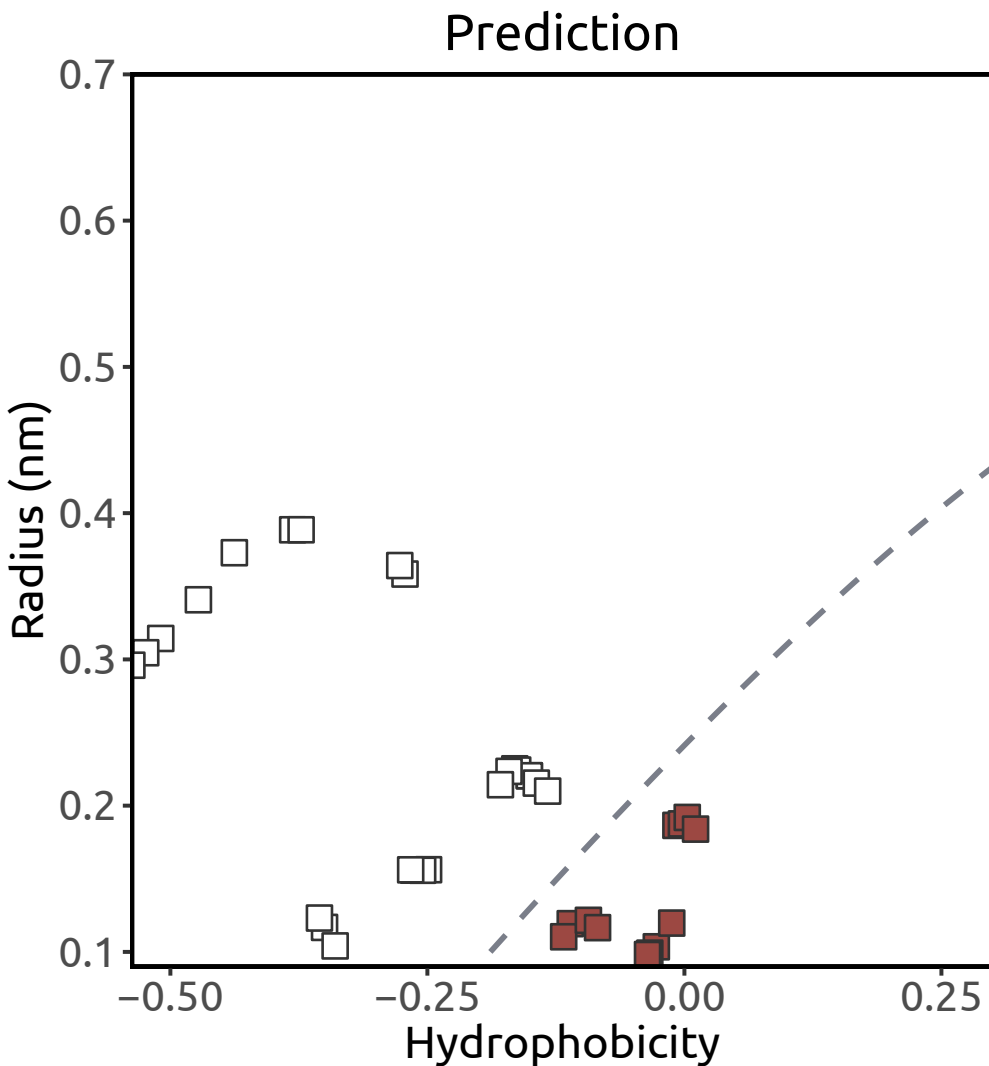
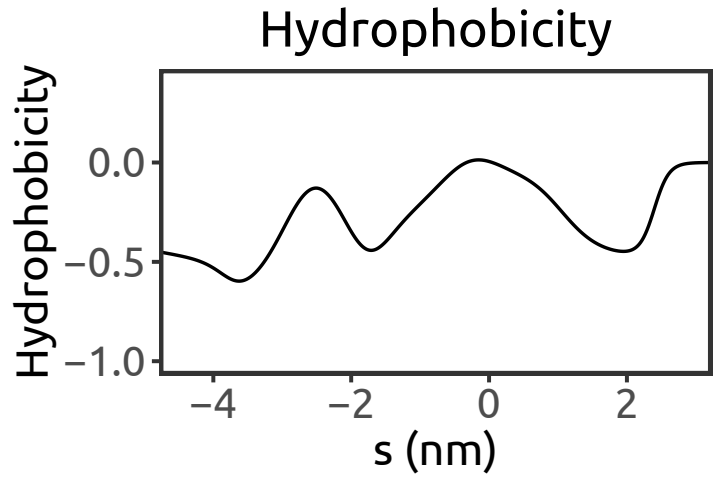
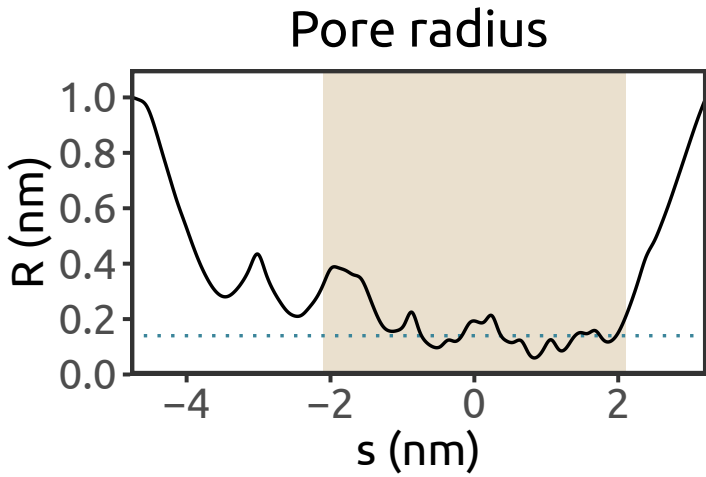
Heuristic score:
0 (n = 1)

Simulation result:
hydrated channel

CNG (PDB ID: 5V4S)

Leptospira licerasiae
cryo-EM (4.2 Å)

James et al., 2017



Heuristic score:
0.71 (n = 12)

Simulation result:
barrier to water

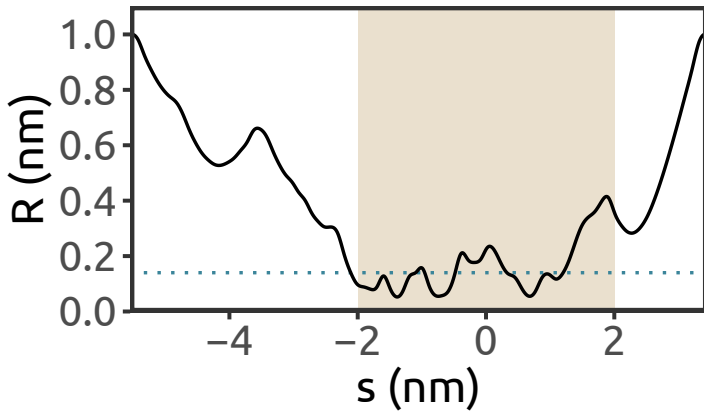
HCN1 (PDB ID: 5U6O)

Homo sapiens

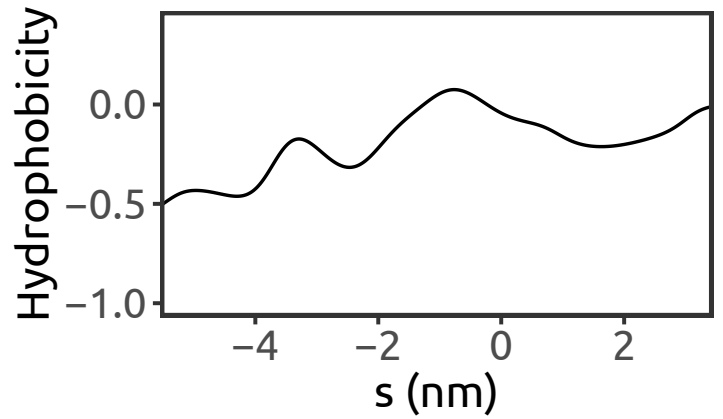
cryo-EM (3.5 Å)

Lee & MacKinnon, 2017

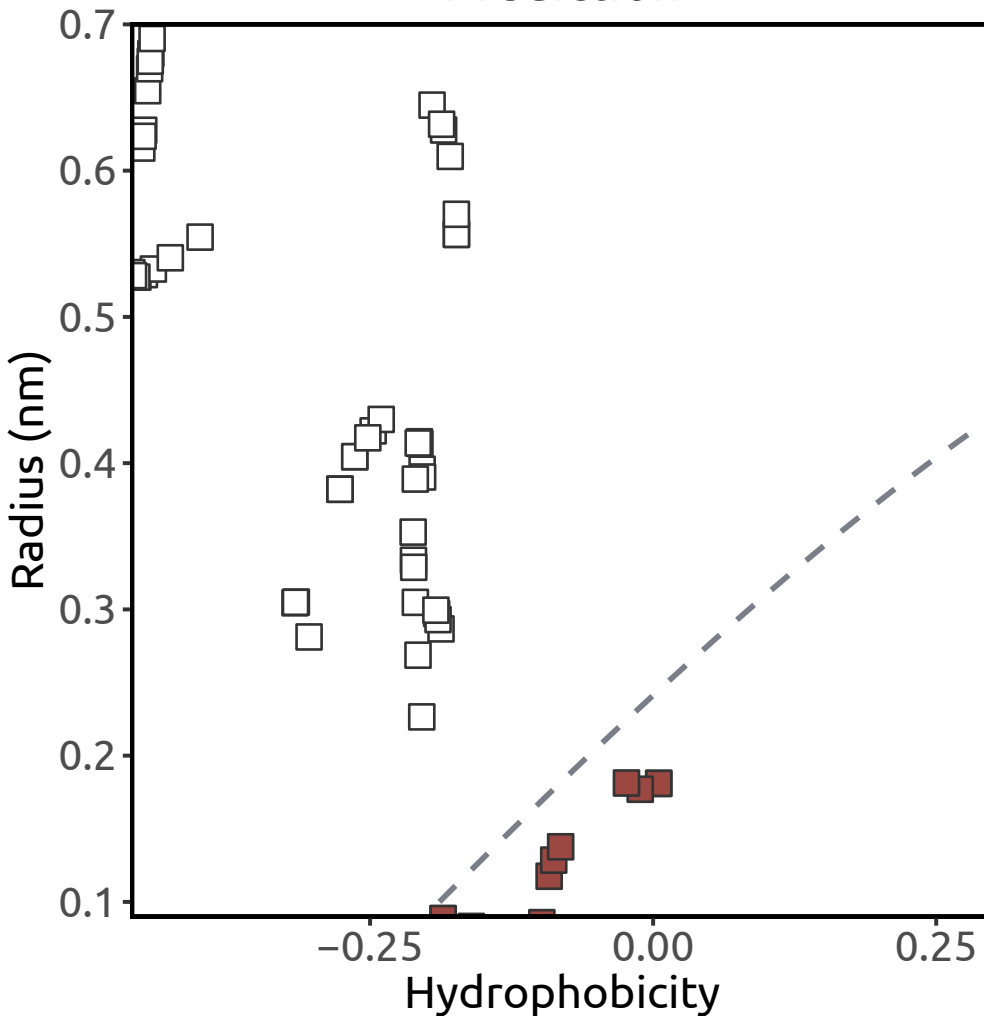
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.78 (n = 20)

Simulation result:

barrier to water

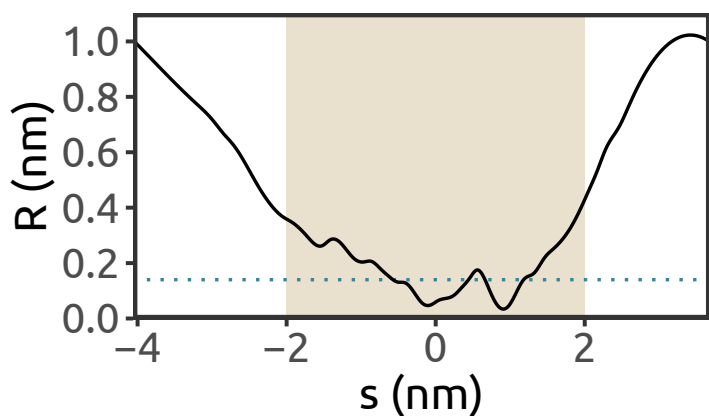
Hv1 (PDB ID: 3WKV)

Mus musculus

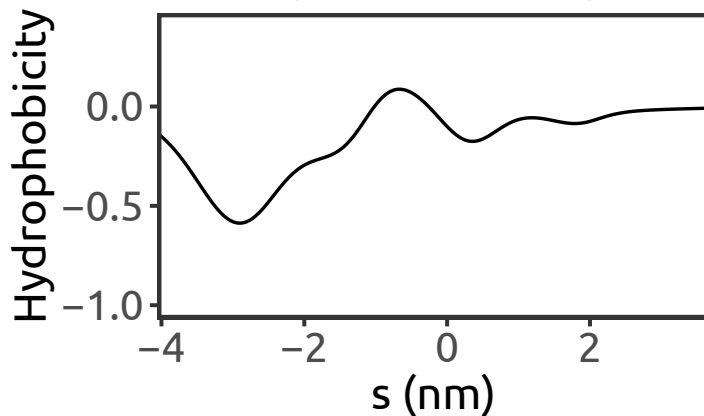
X-ray (3.45 Å)

Takeshita et al., 2014

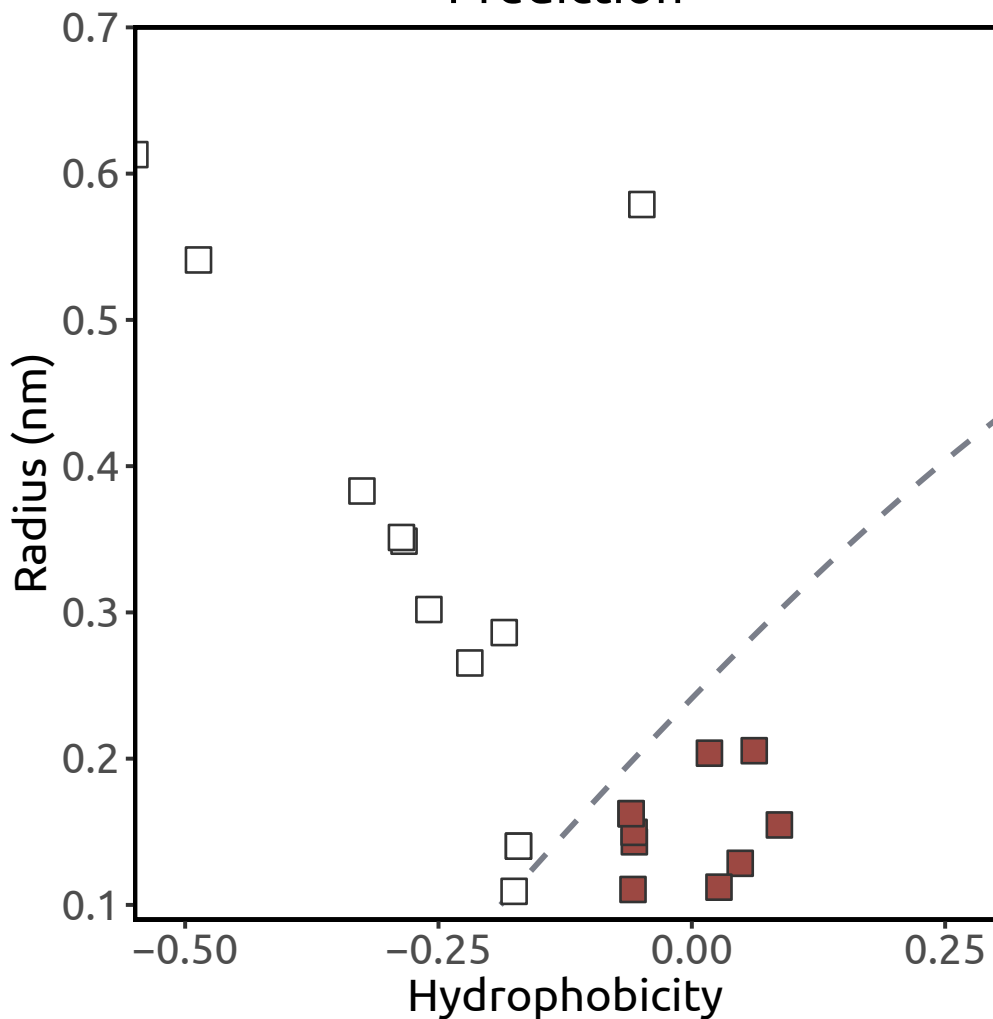
Pore radius



Hydrophobicity



Prediction



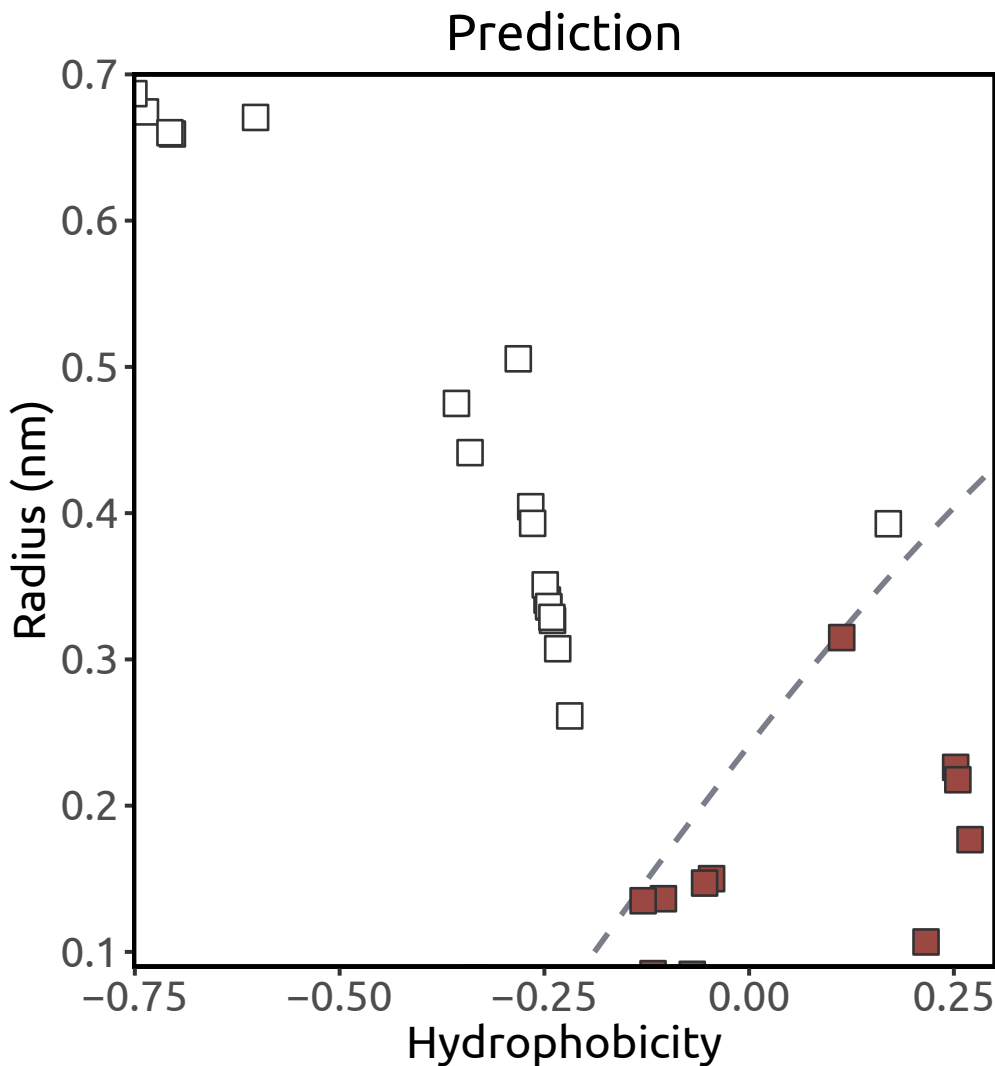
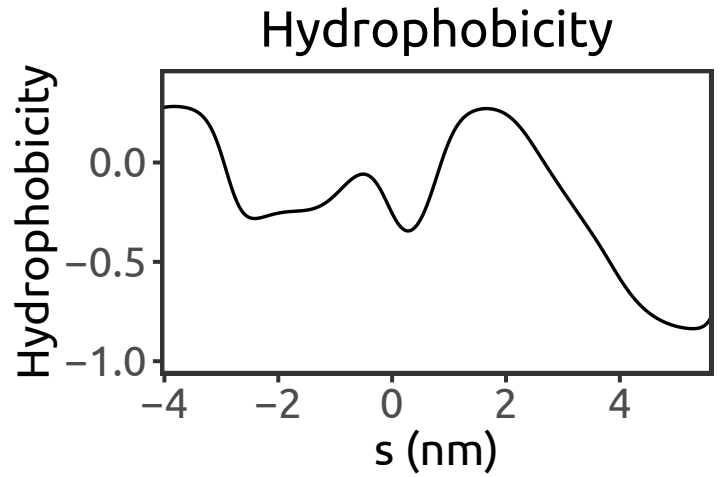
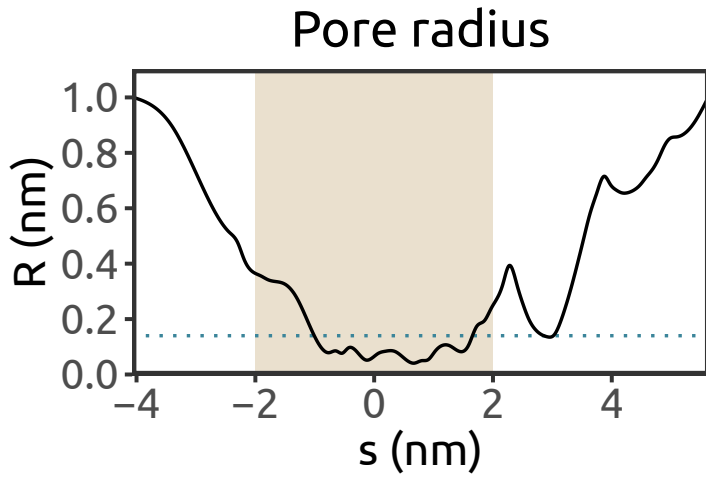
Heuristic score:
1.35 (n = 17)

Simulation result:
barrier to water

InsP3R1 (PDB ID: 3JAV)

Rattus norvegicus
cryo-EM (4.7 Å)

Fan et al., 2015



Heuristic score:
1.02 (n = 11)

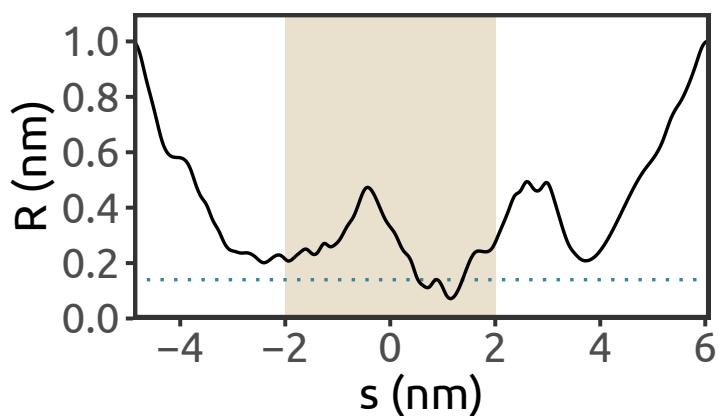
Simulation result:
barrier to water

Nav1.4 (PDB ID: 5XSY)

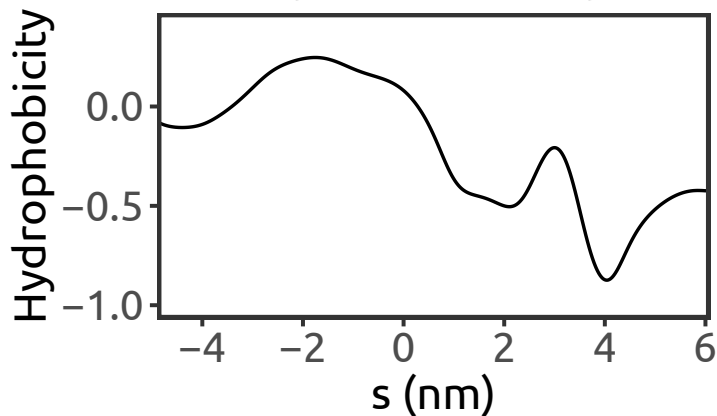
Electrophorus electricus
cryo-EM (4 Å)

Yan et al., 2017

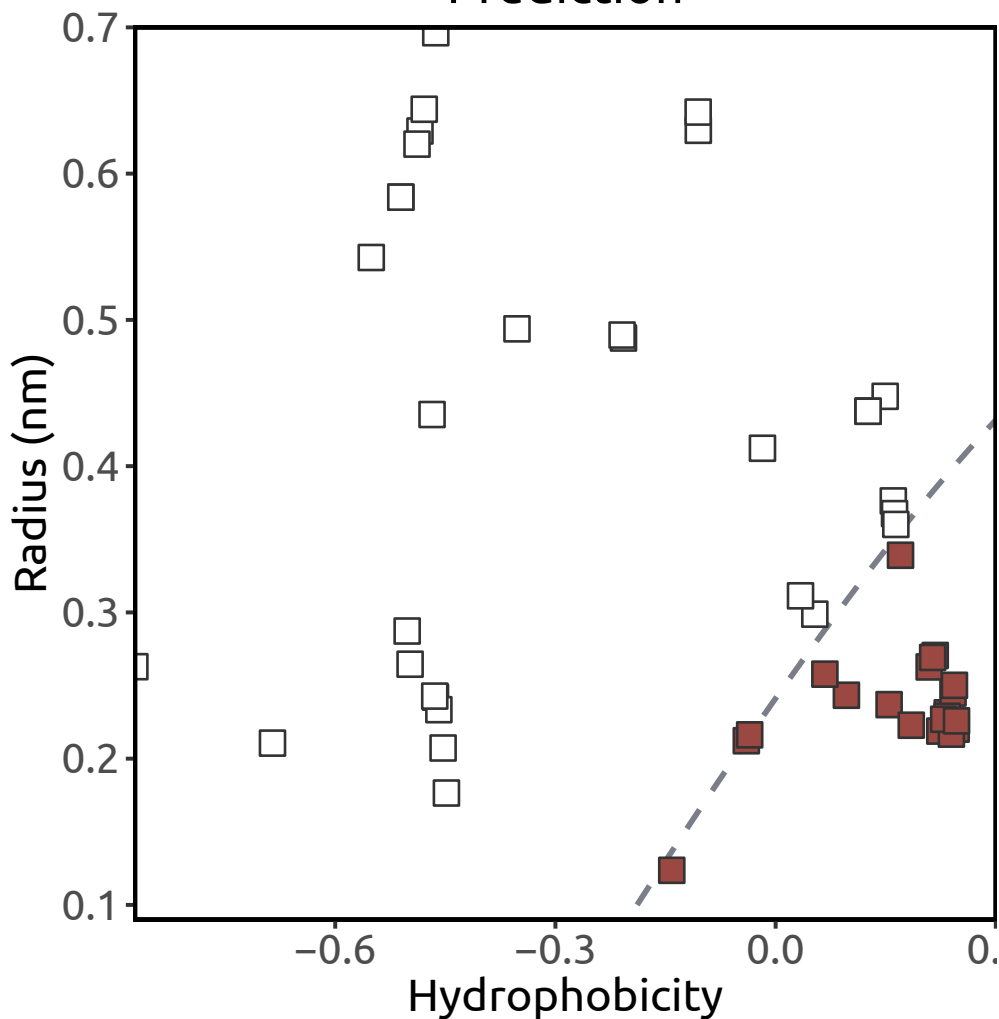
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.04 (n = 21)

Simulation result:
barrier to water

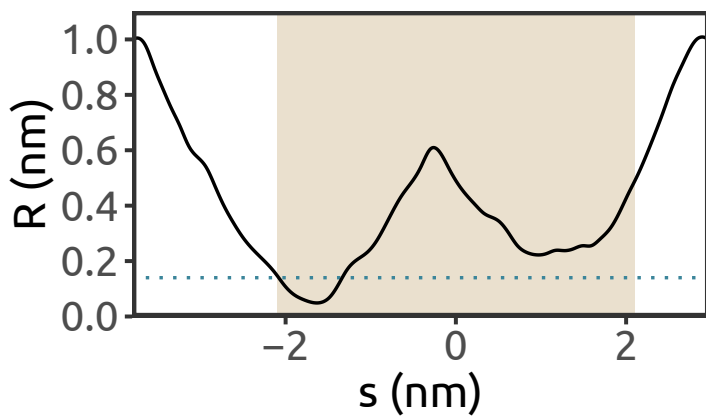
NavAb (PDB ID: 4EKW)

Arcobacter butzleri

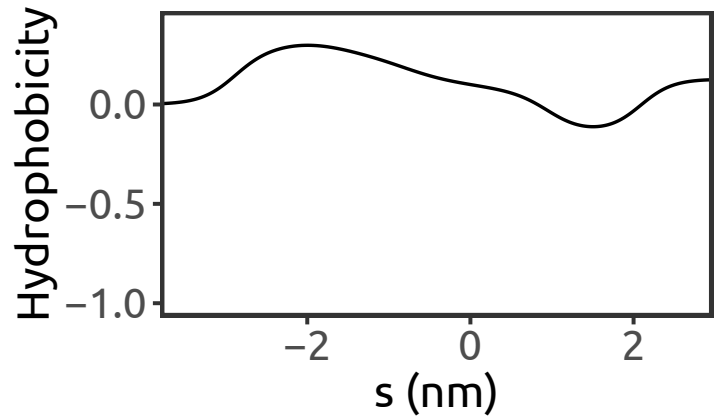
X-ray (3.21 Å)

Payandeh et al., 2012

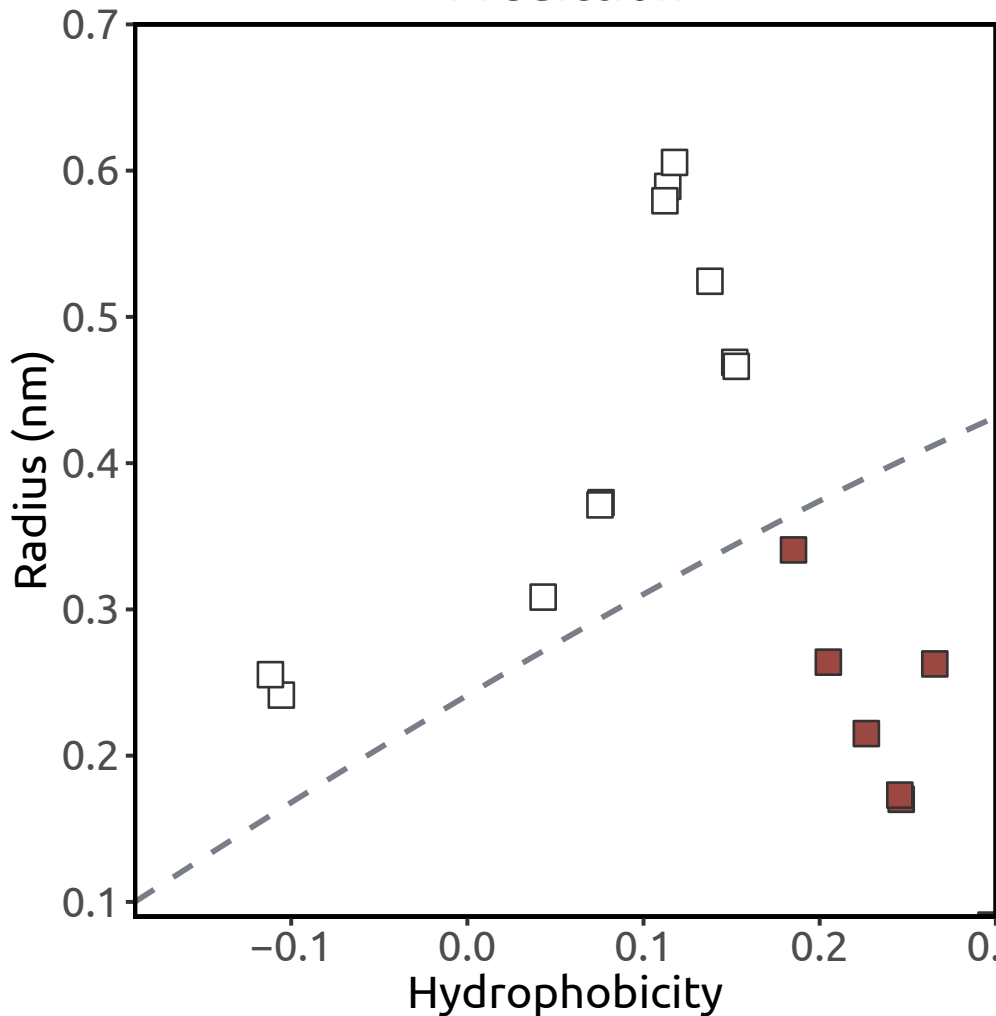
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.72 (n = 9)

Simulation result:
barrier to water

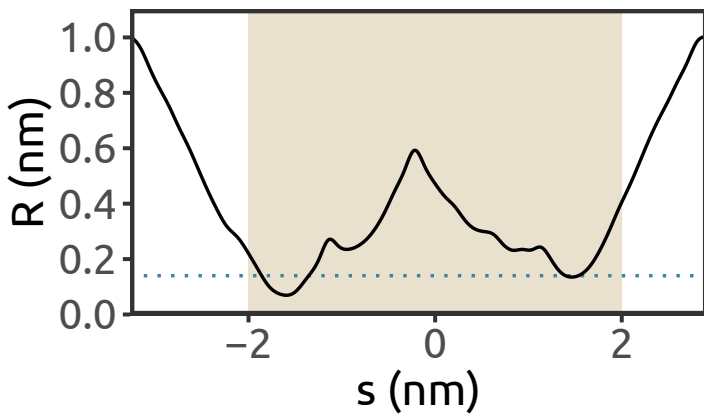
NavAb (PDB ID: 4MW8)

Arcobacter butzleri

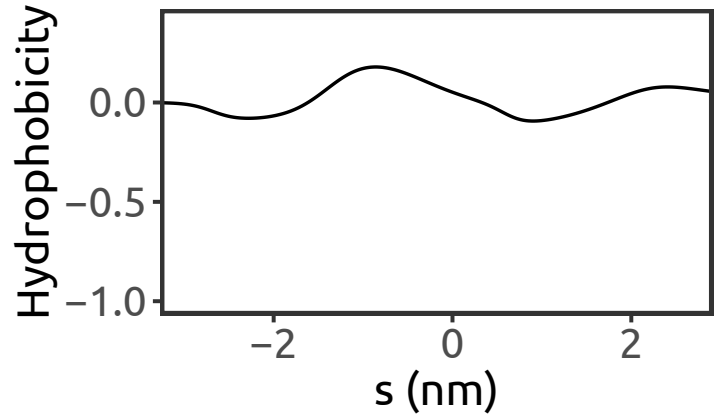
X-ray (3.26 Å)

Tang et al., 2014

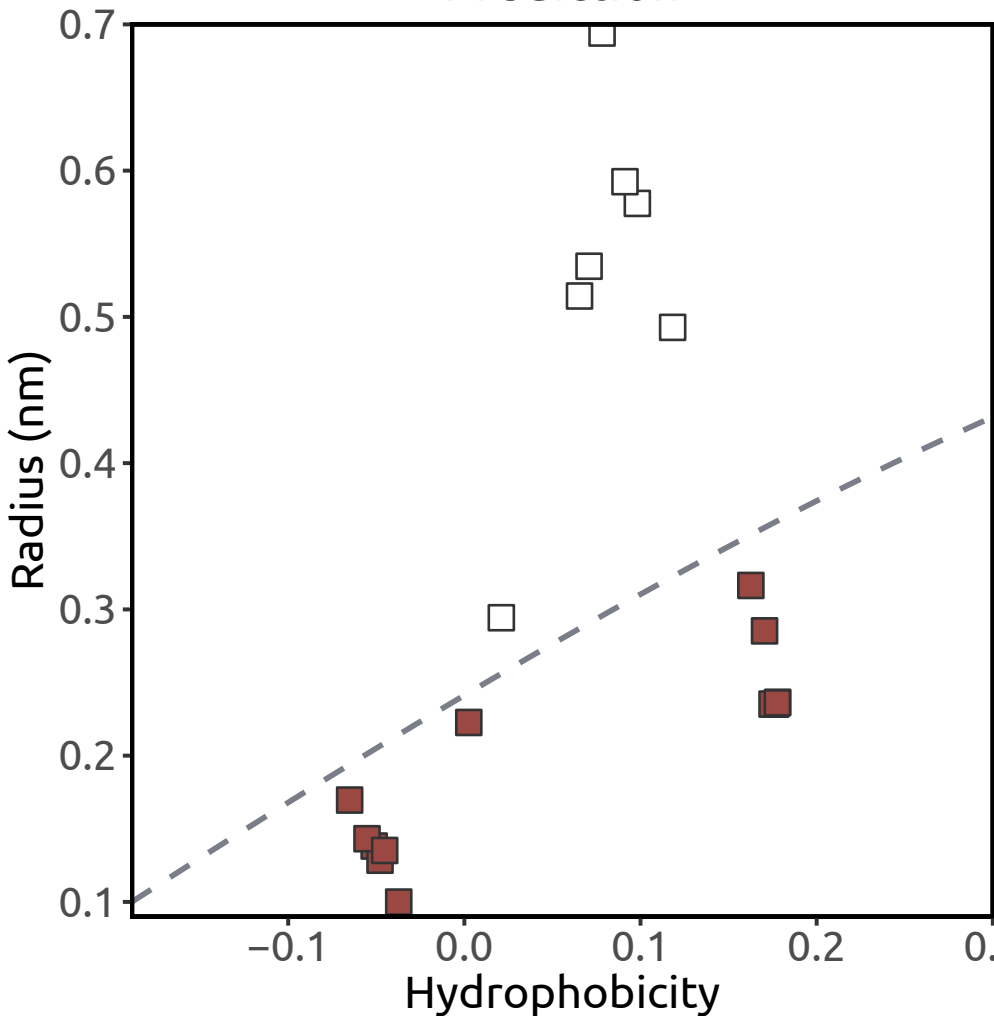
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.11 (n = 15)

Simulation result:
hydrated channel

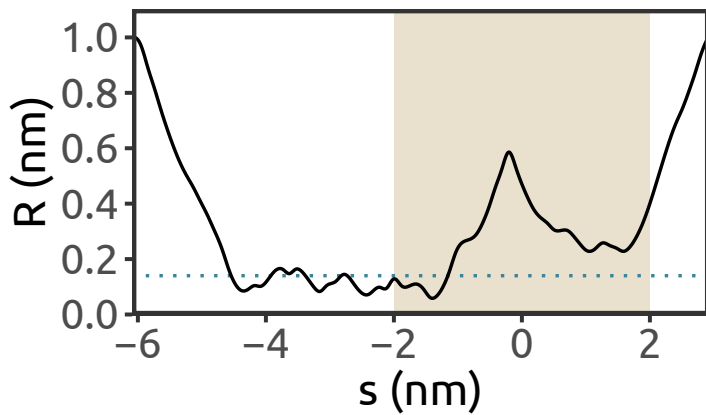
NavAb (PDB ID: 5EK0)

Arcobacter butzleri

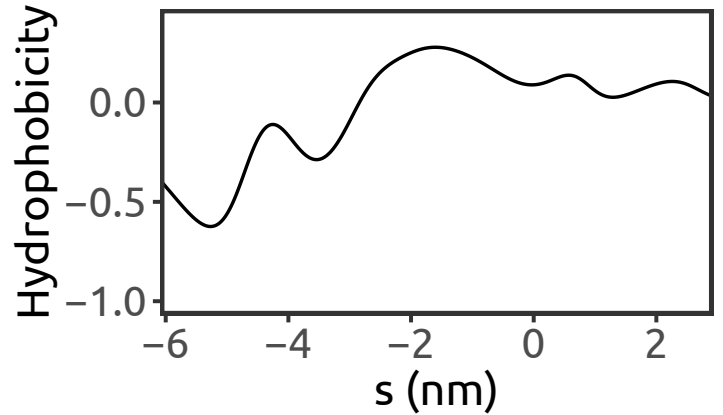
X-ray (3.53 Å)

Ahuja et al., 2015

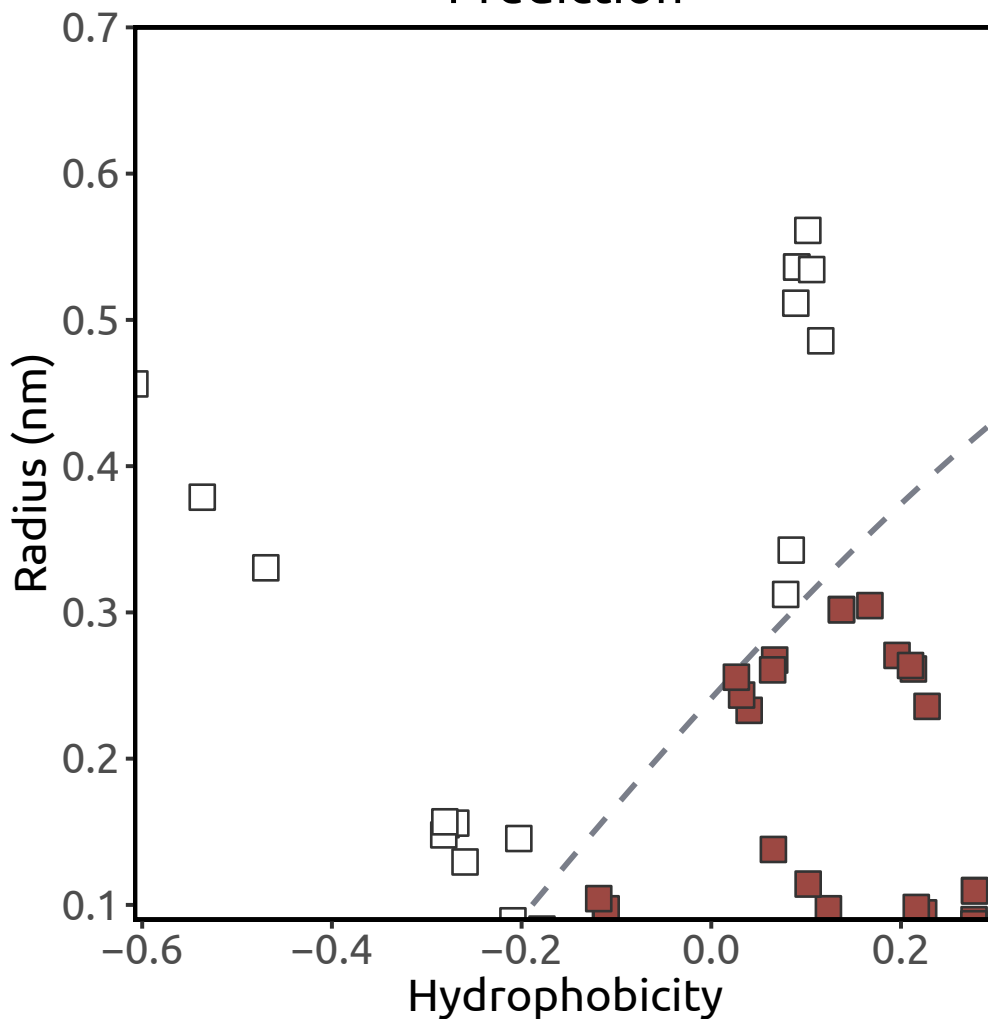
Pore radius



Hydrophobicity



Prediction



Heuristic score:
3.4 (n = 28)

Simulation result:
barrier to water

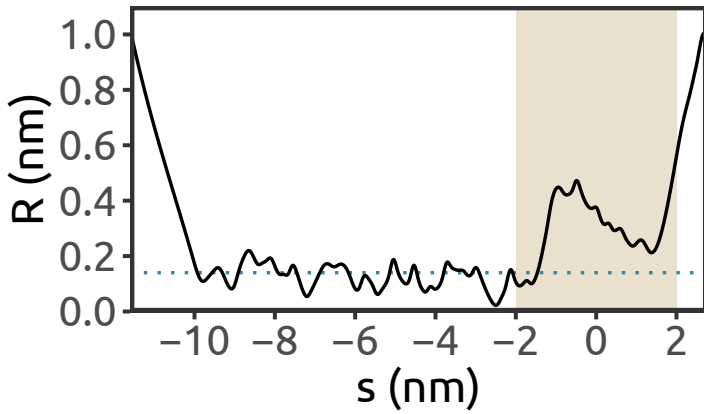
NavAb (PDB ID: 5VB2)

Arcobacter butzleri

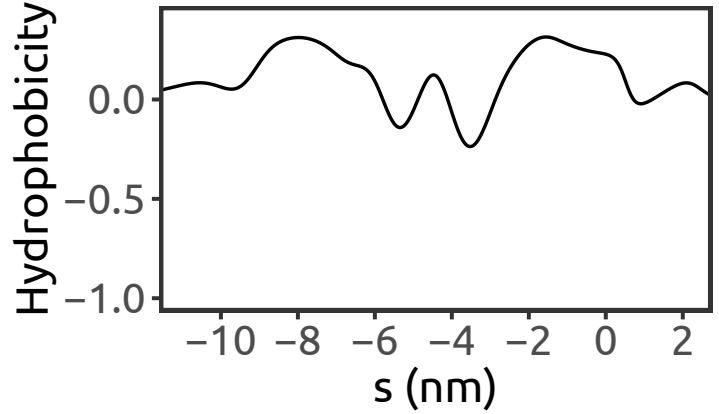
X-ray (3.2 Å)

Lenaeus et al., 2017

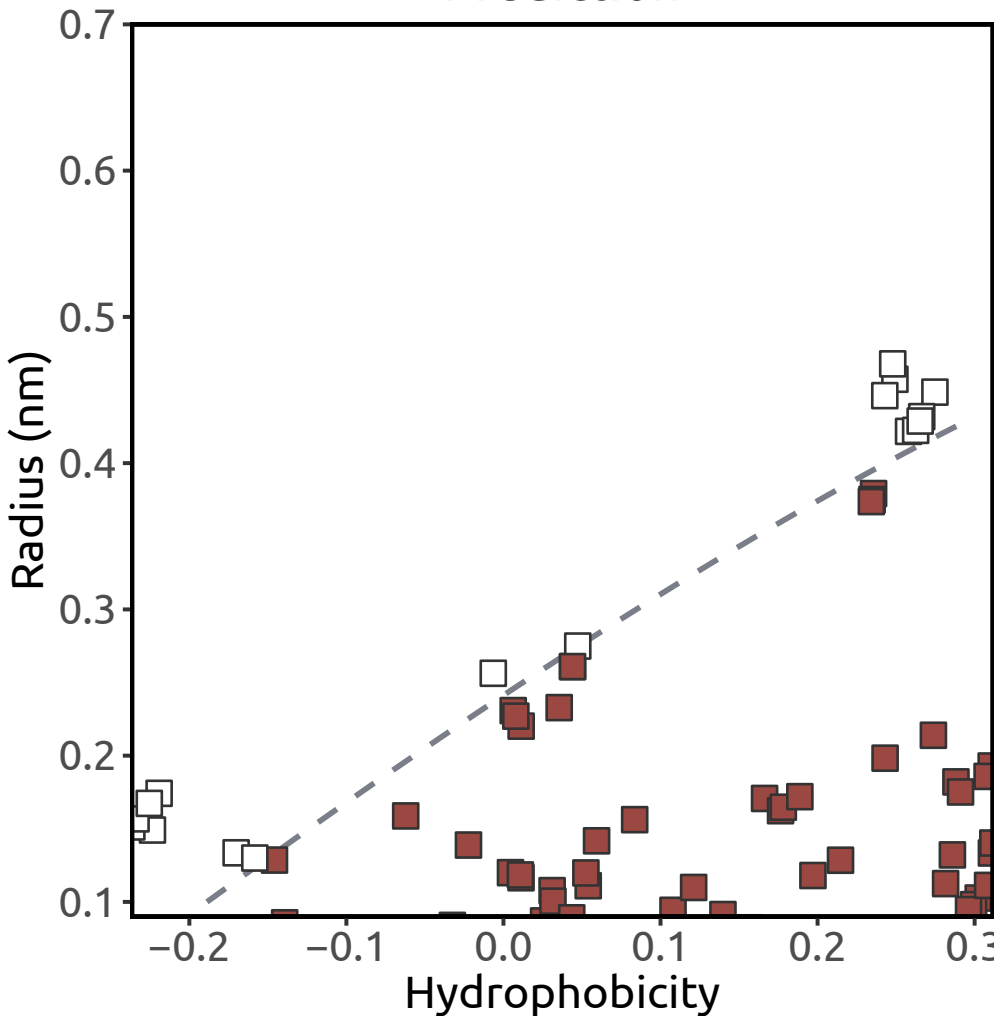
Pore radius



Hydrophobicity



Prediction



Heuristic score:

9.06 (n = 61)

Simulation result:

barrier to water

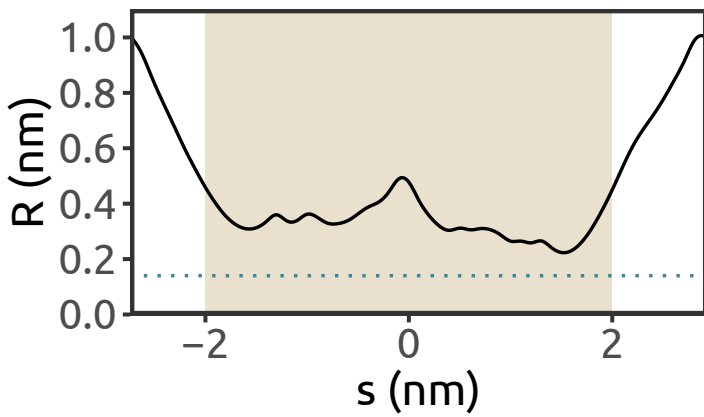
NavAb (PDB ID: 5VB8)

Arcobacter butzleri

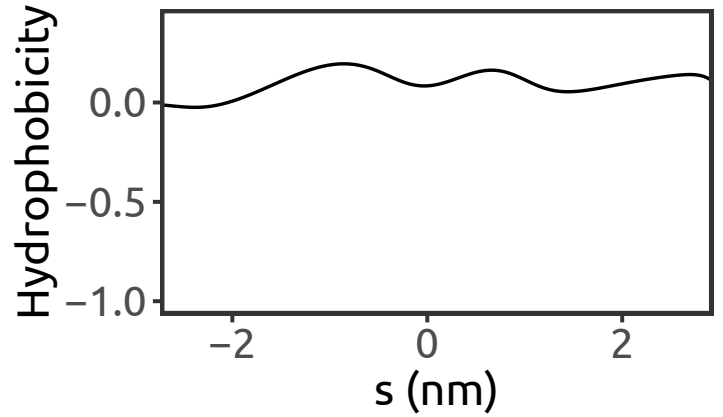
X-ray (2.85 Å)

Lenaeus et al., 2017

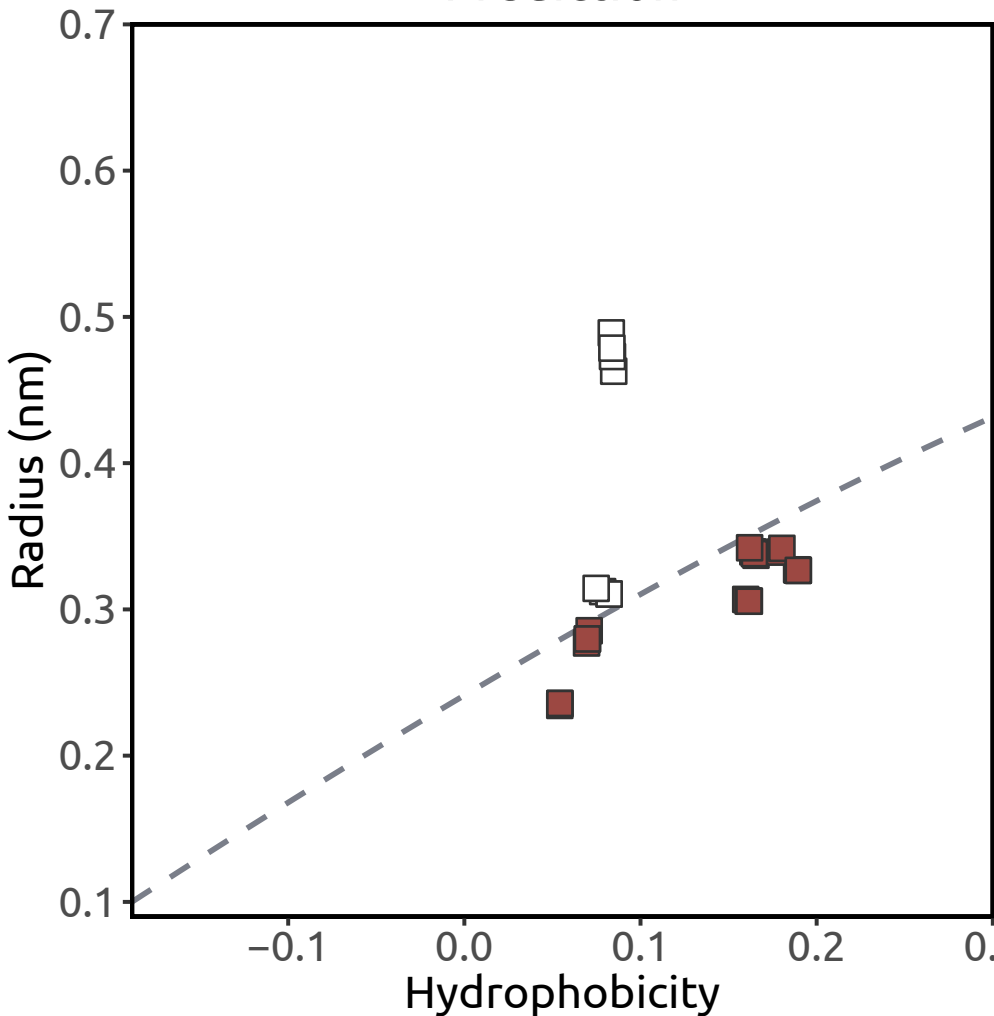
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.51 (n = 21)

Simulation result:
barrier to water

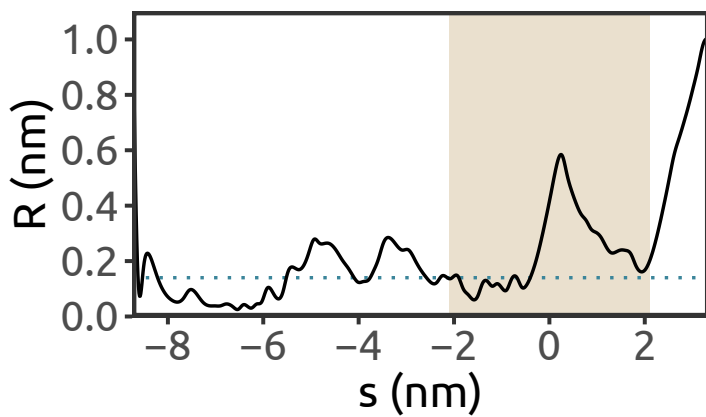
NavAe1 (PDB ID: 4LTO)

Alkalilimnicola ehrlichii

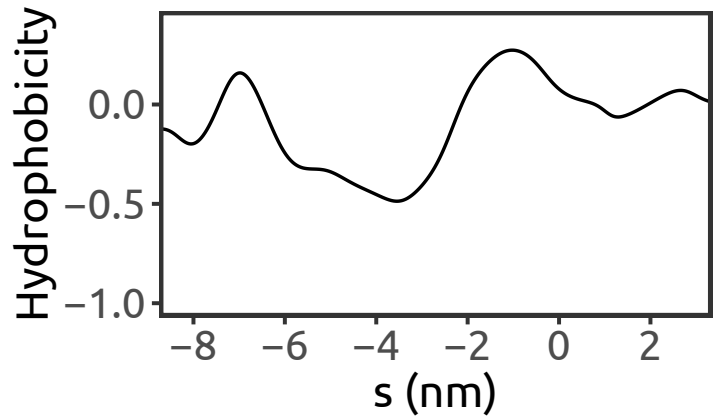
X-ray (3.46 Å)

Shaya et al., 2014

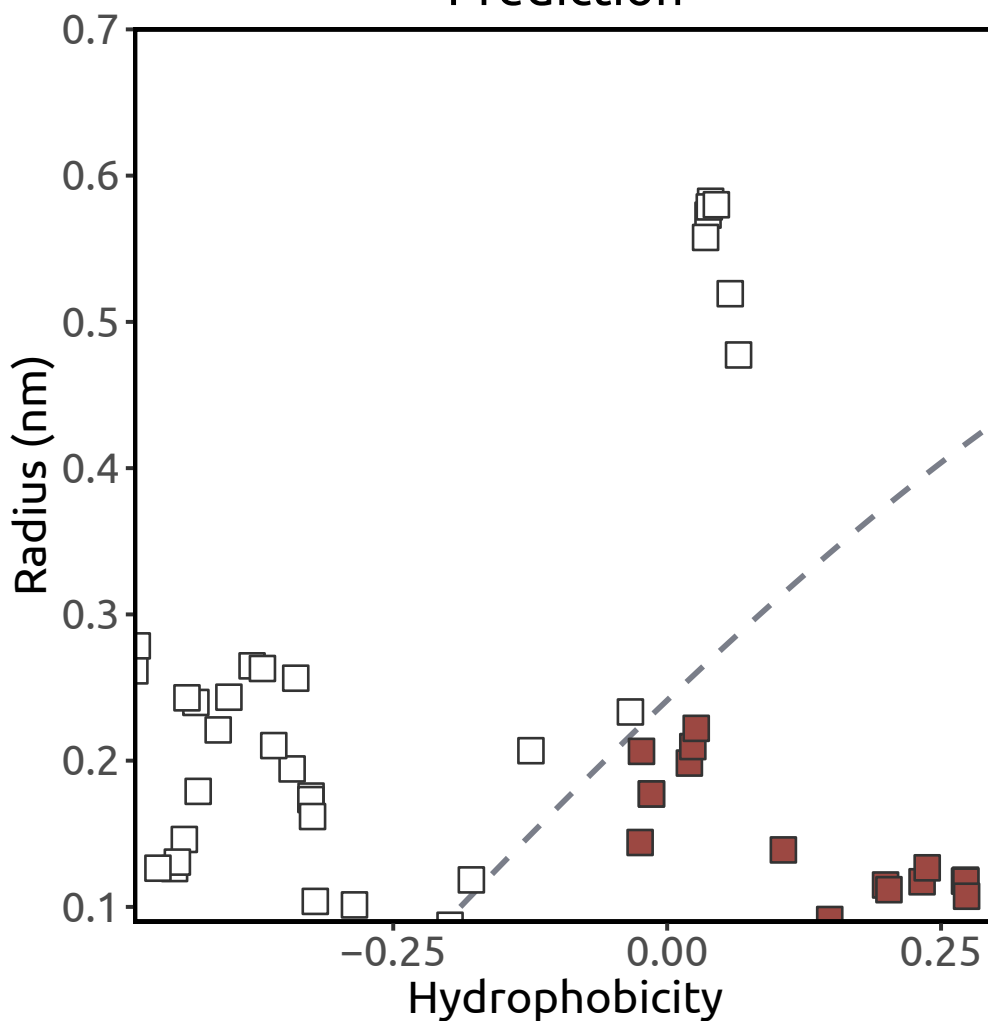
Pore radius



Hydrophobicity



Prediction



Heuristic score:
3.76 (n = 24)

Simulation result:
barrier to water

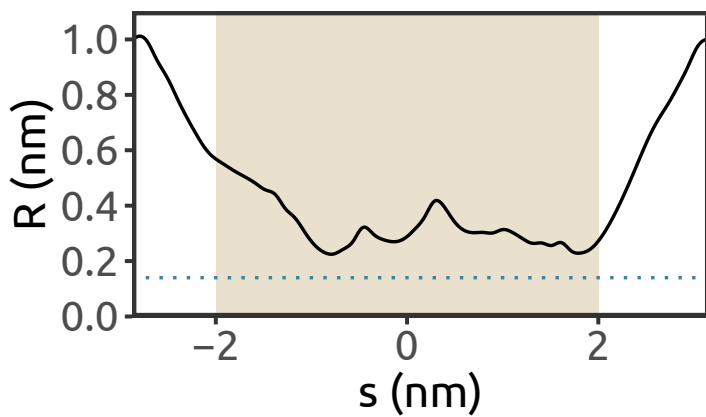
NavMm (PDB ID: 4CBC)

Magnetococcus marinus

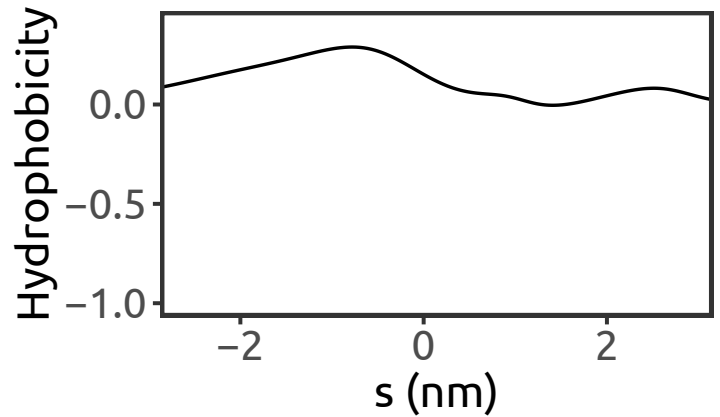
X-ray (2.66 Å)

Bagn ris et al., 2014

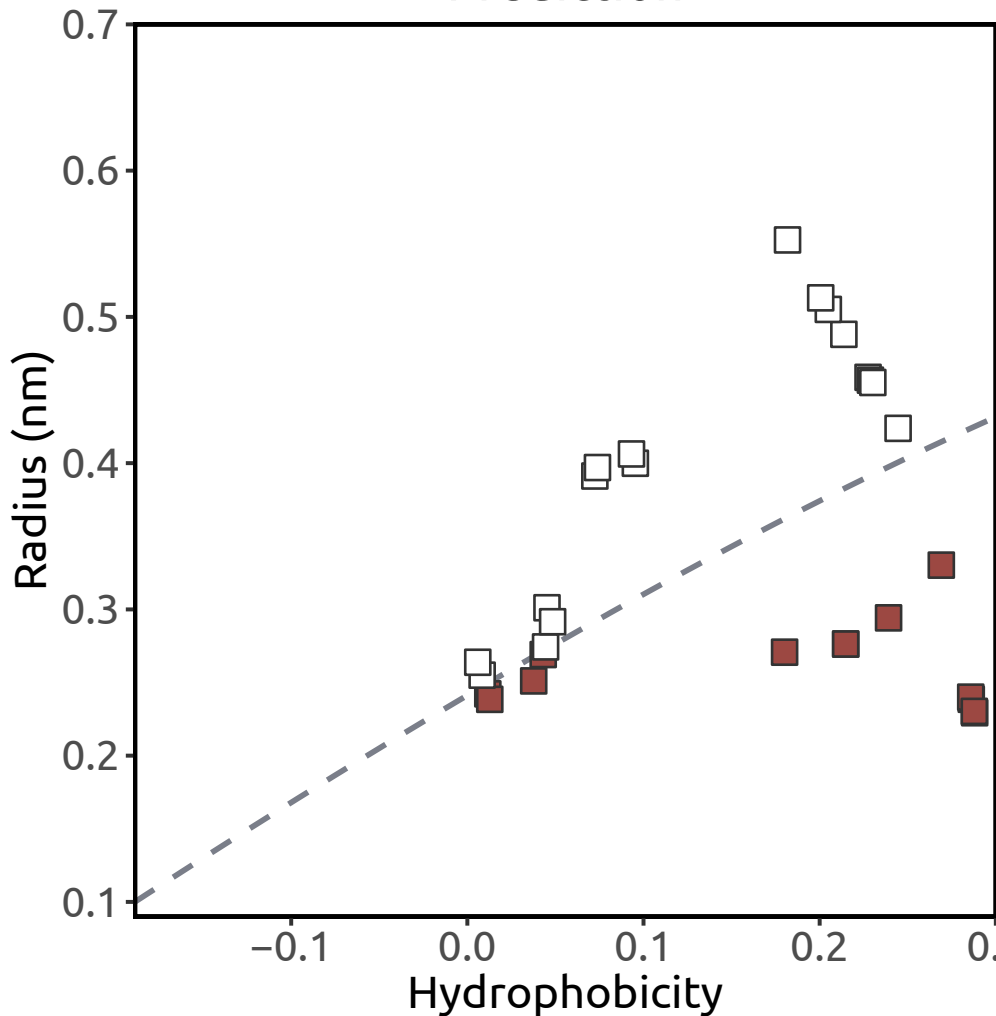
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.11 (n = 13)

Simulation result:
barrier to water

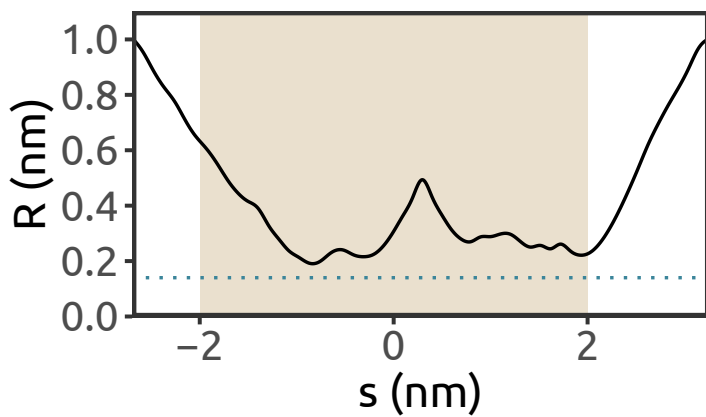
NavMm (PDB ID: 4OXS)

Magnetococcus marinus

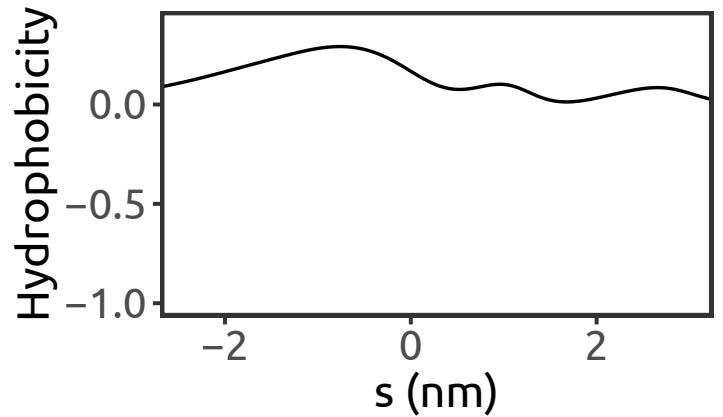
X-ray (2.8 Å)

Bagn ris et al., 2014

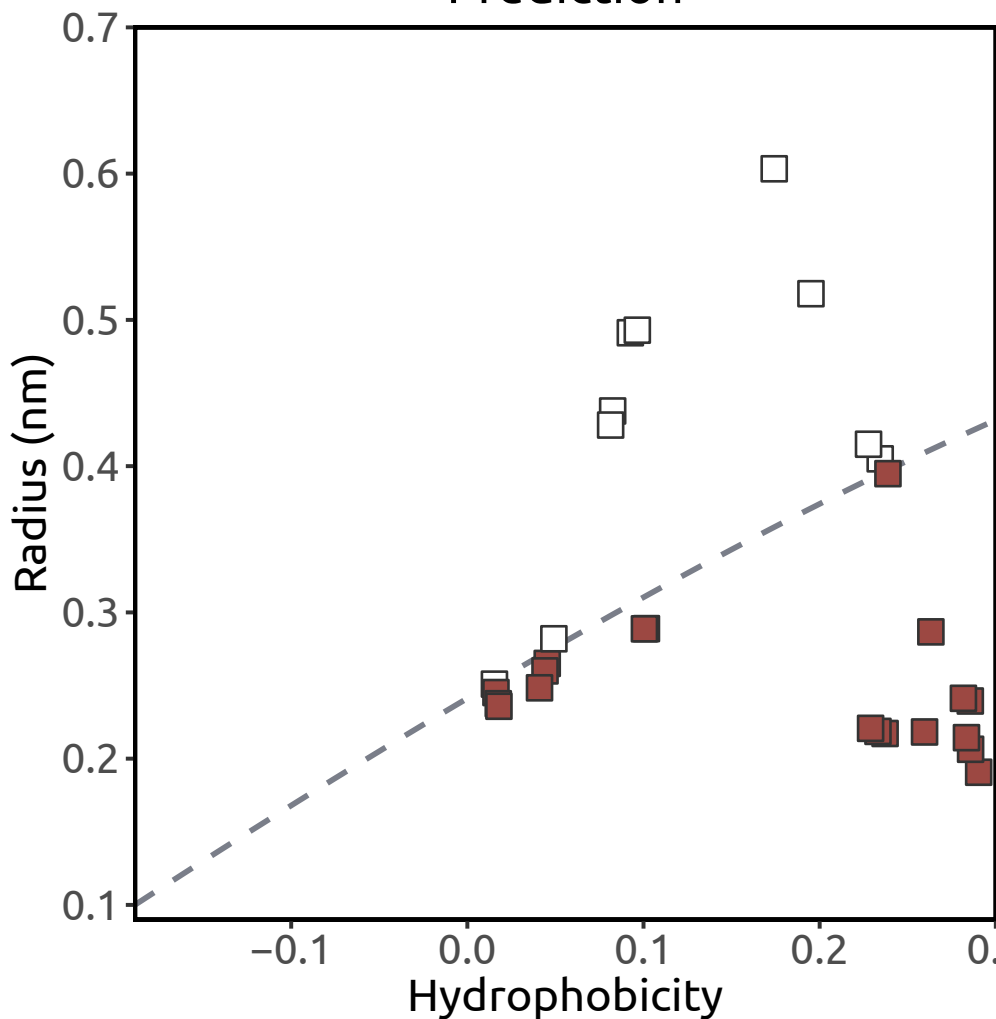
Pore radius



Hydrophobicity



Prediction



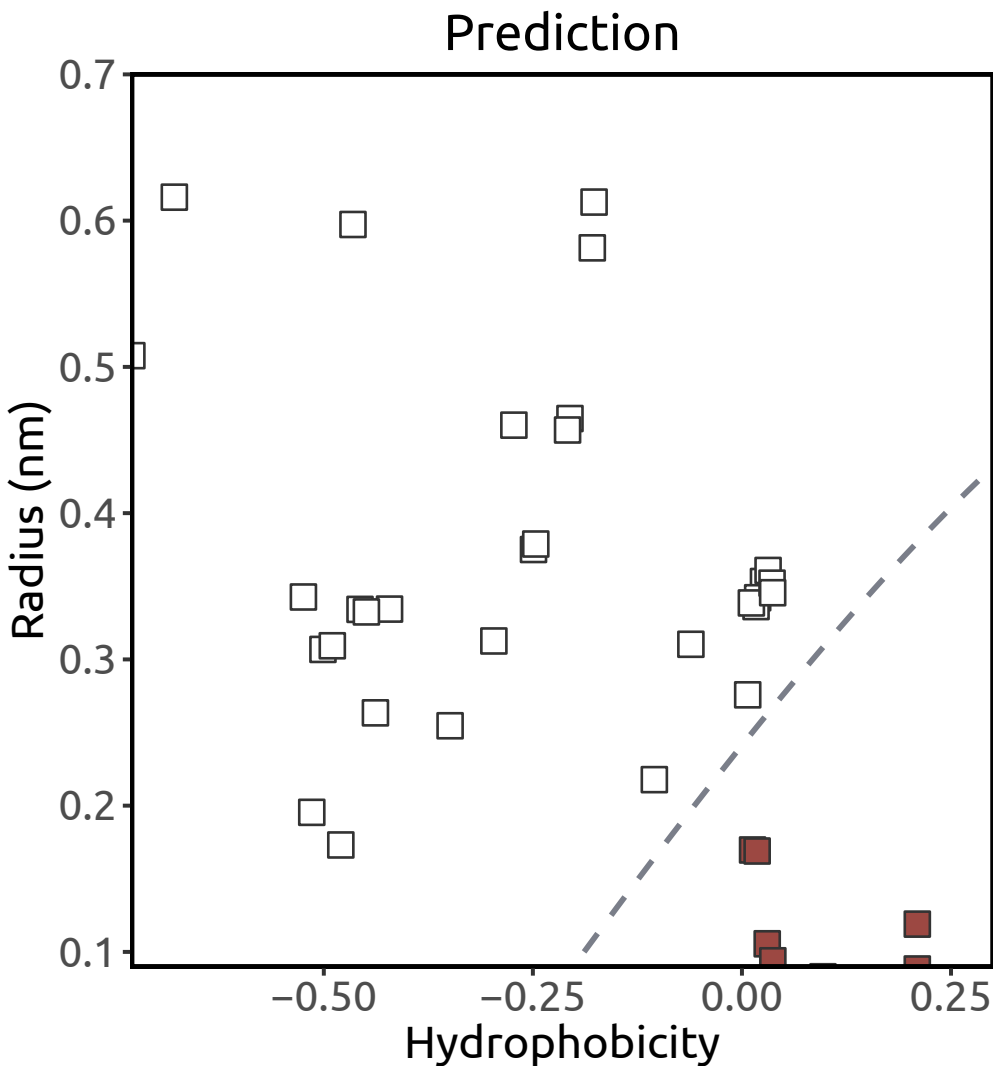
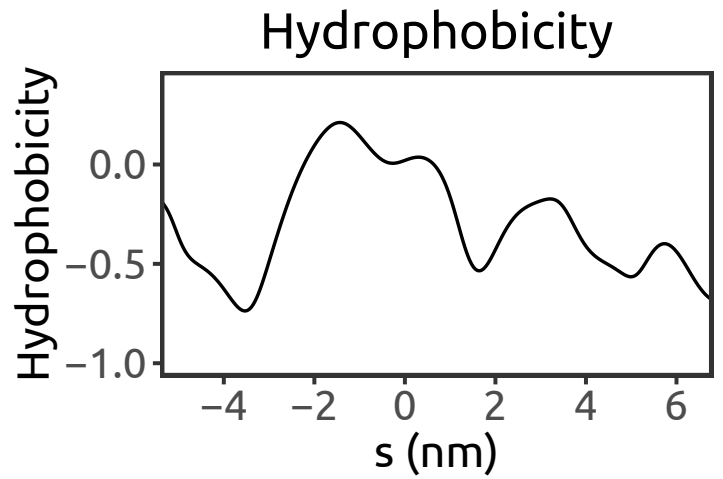
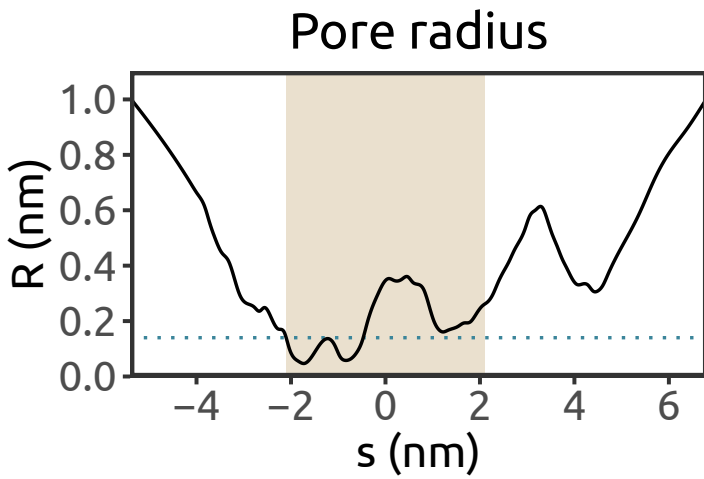
Heuristic score:
1.72 (n = 19)

Simulation result:
barrier to water

NavPa (PDB ID: 5X0M)

Periplaneta americana
cryo-EM (3.8 Å)

Shen et al., 2017



Heuristic score:
1.91 (n = 11)

Simulation result:
barrier to water

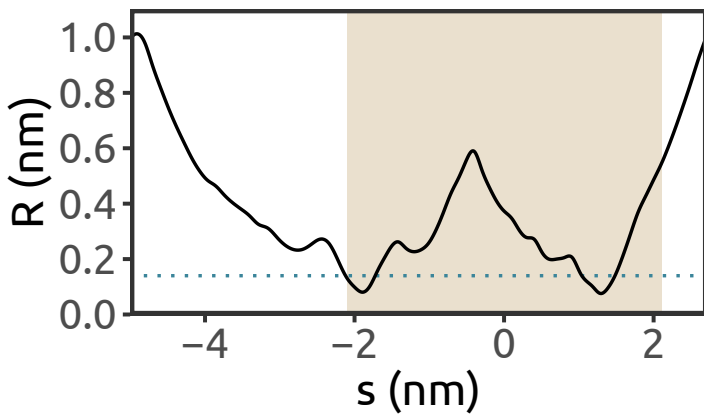
NavRh (PDB ID: 4DXW)

Rickettsiales sp. HIMB114

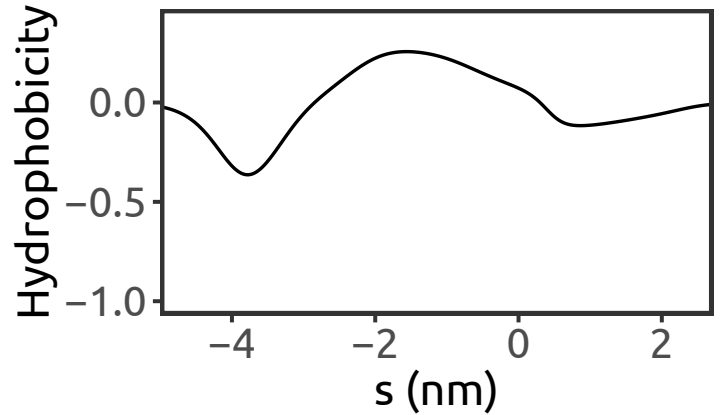
X-ray (3.05 Å)

Zhang et al., 2012

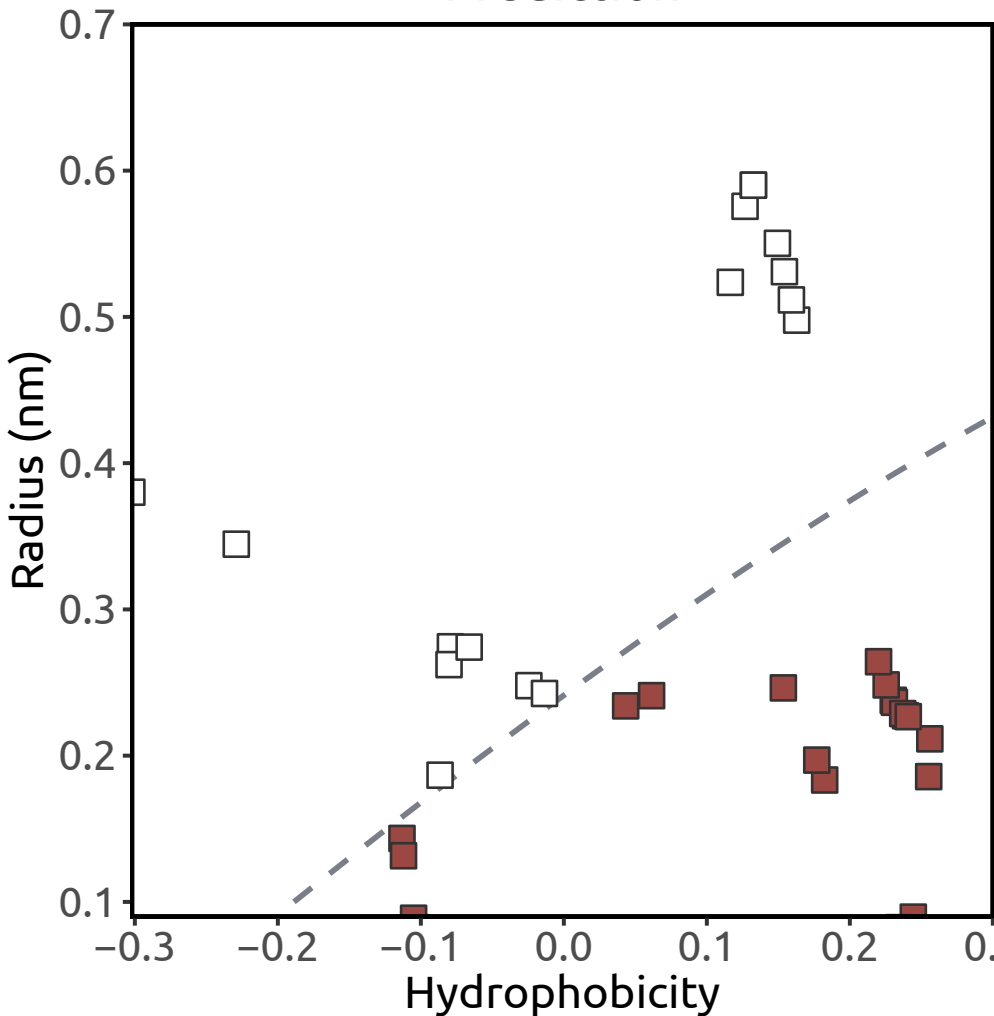
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.67 (n = 21)

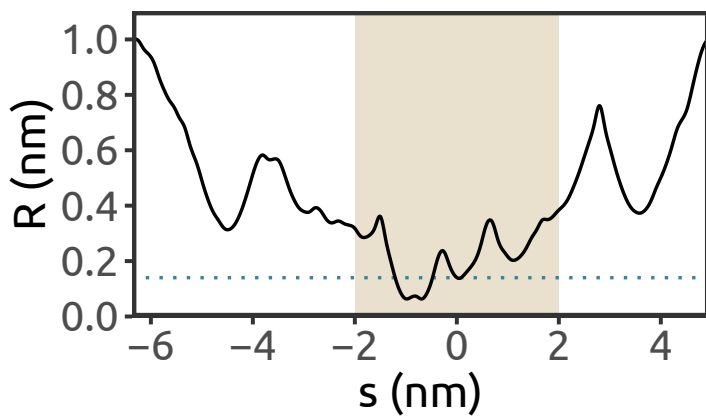
Simulation result:
barrier to water

RyR1 (PDB ID: 3J8H)

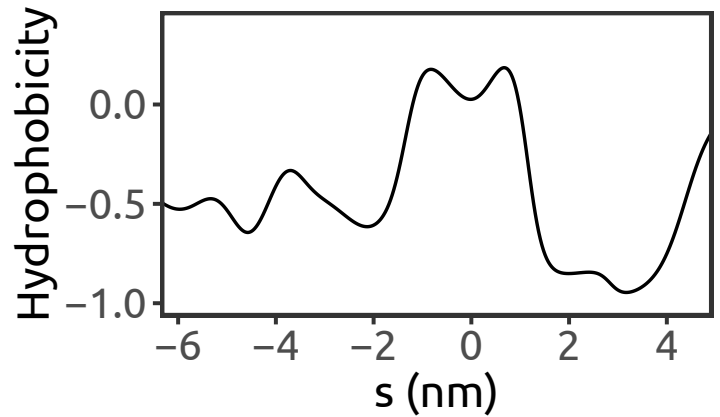
Oryctolagus cuniculus
cryo-EM (3.8 Å)

Yan et al., 2015

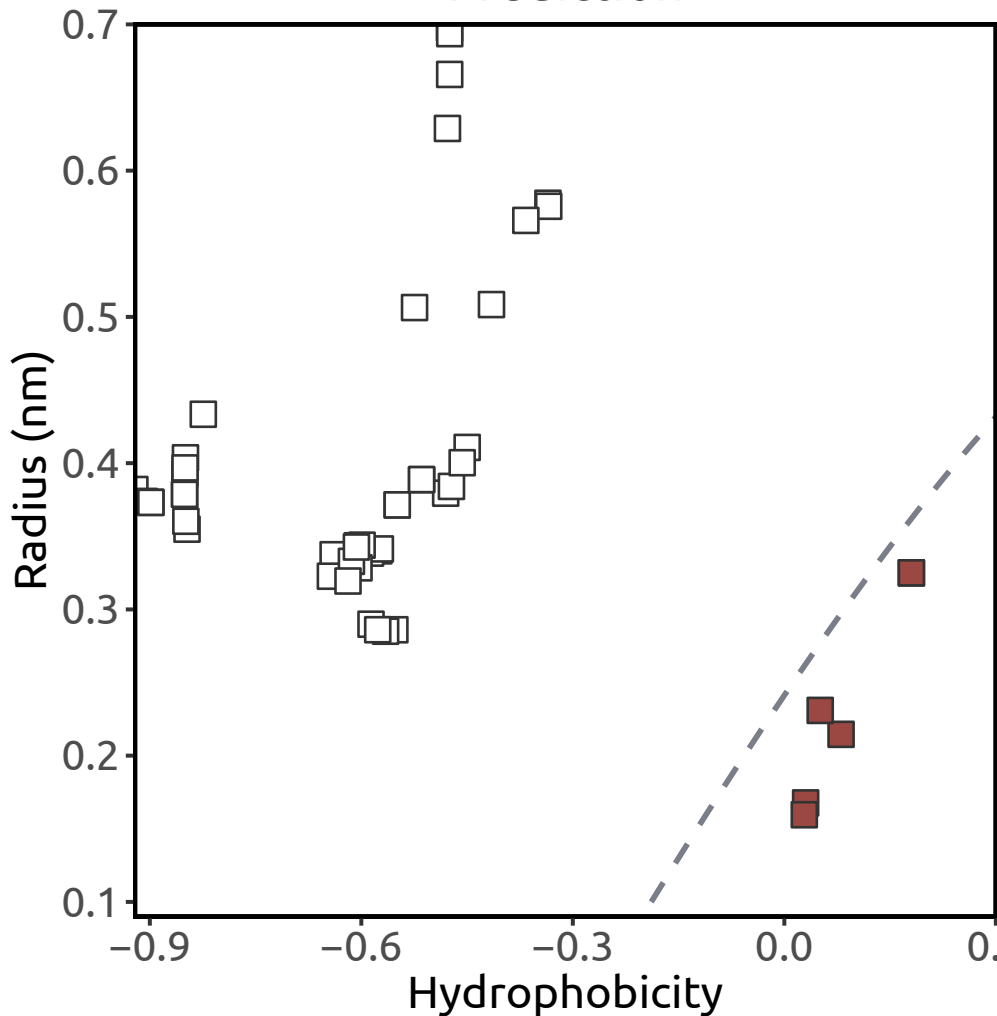
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.29 ($n = 10$)

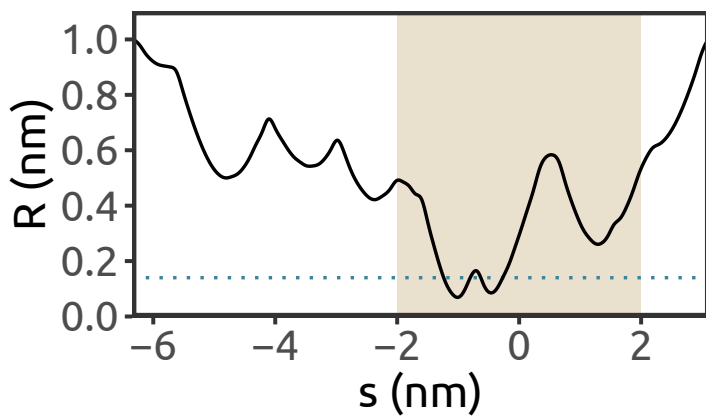
Simulation result:
barrier to water

RyR1 (PDB ID: 5TB0)

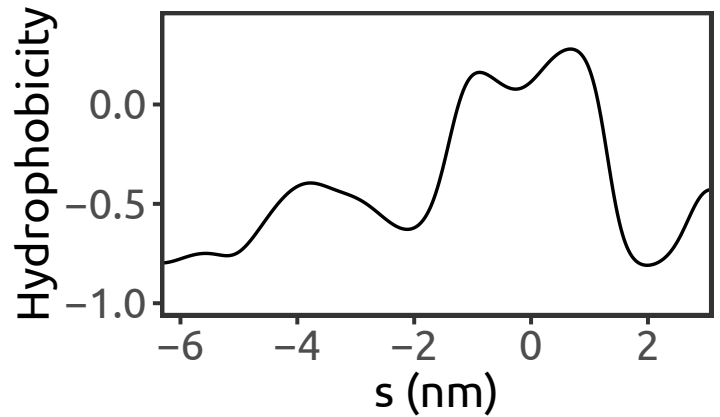
Oryctolagus cuniculus
cryo-EM (4.4 Å)

des et al., 2016

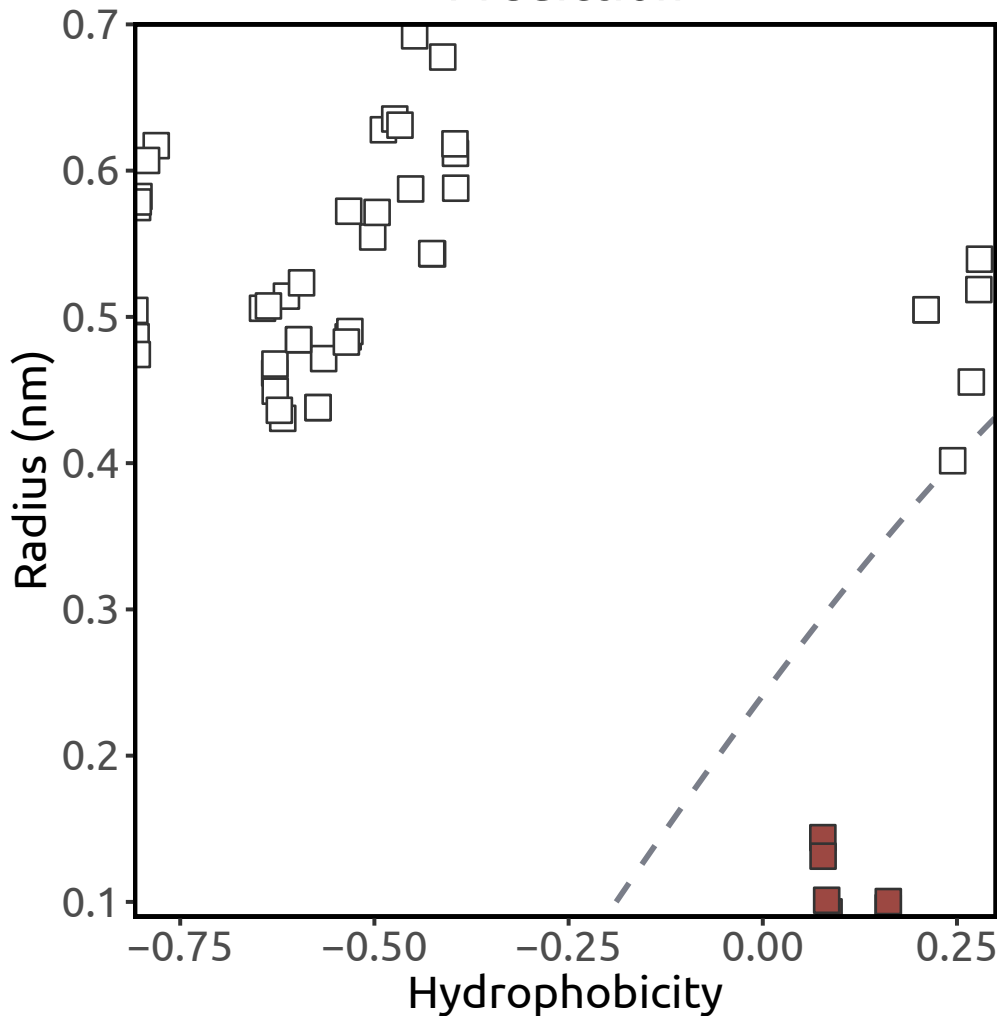
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.46 (n = 8)

Simulation result:
barrier to water

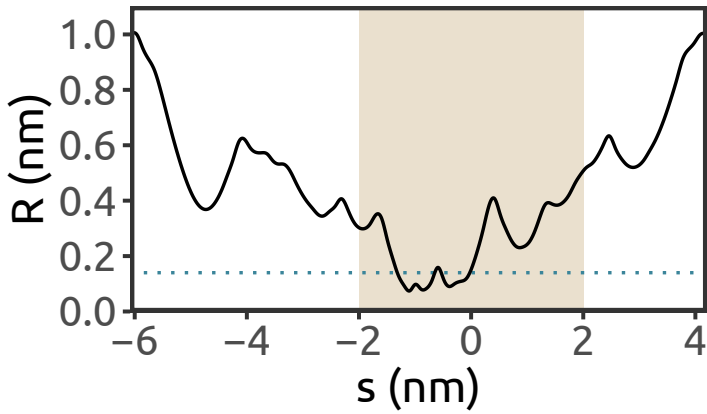
RyR2 (PDB ID: 5GO9)

Sus scrofa

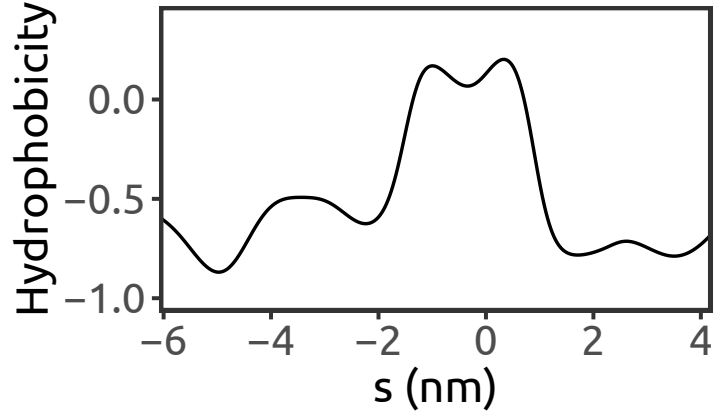
cryo-EM (4.4 Å)

Peng et al., 2016

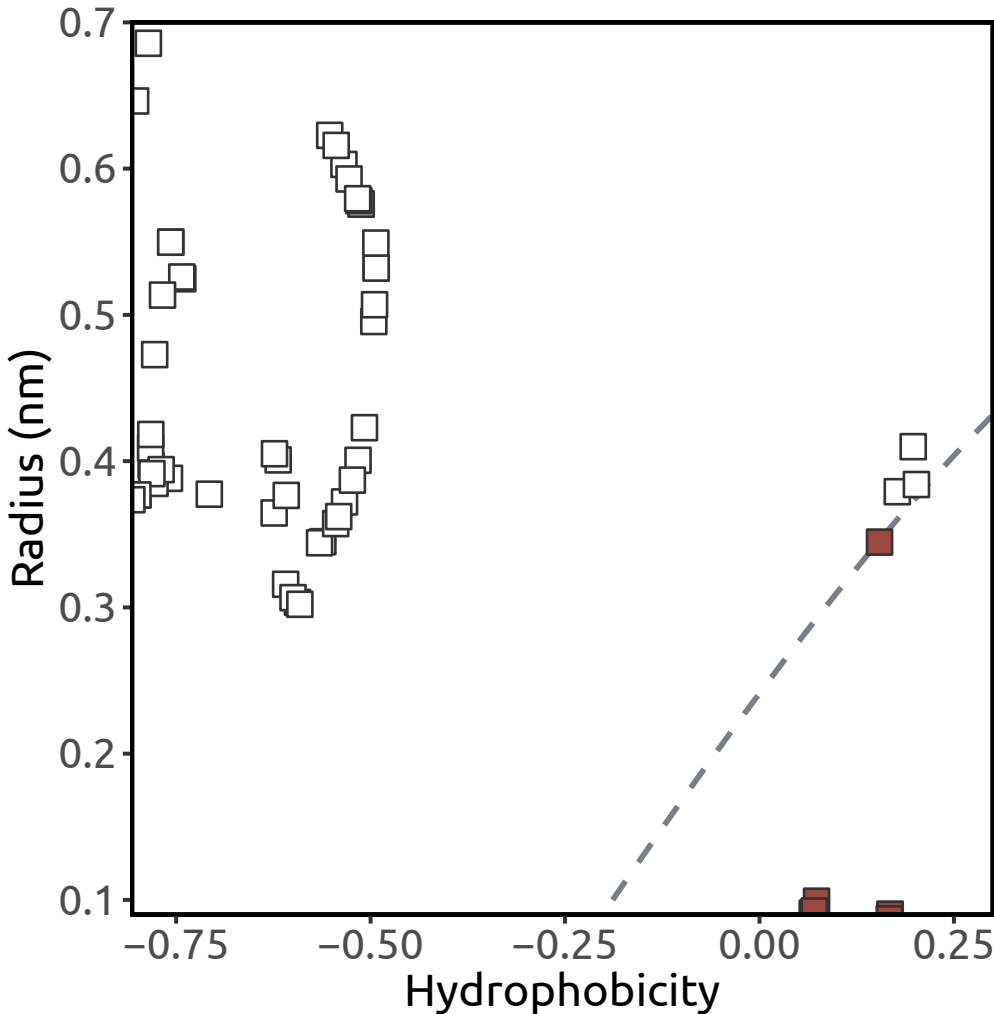
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.53 (n = 9)

Simulation result:

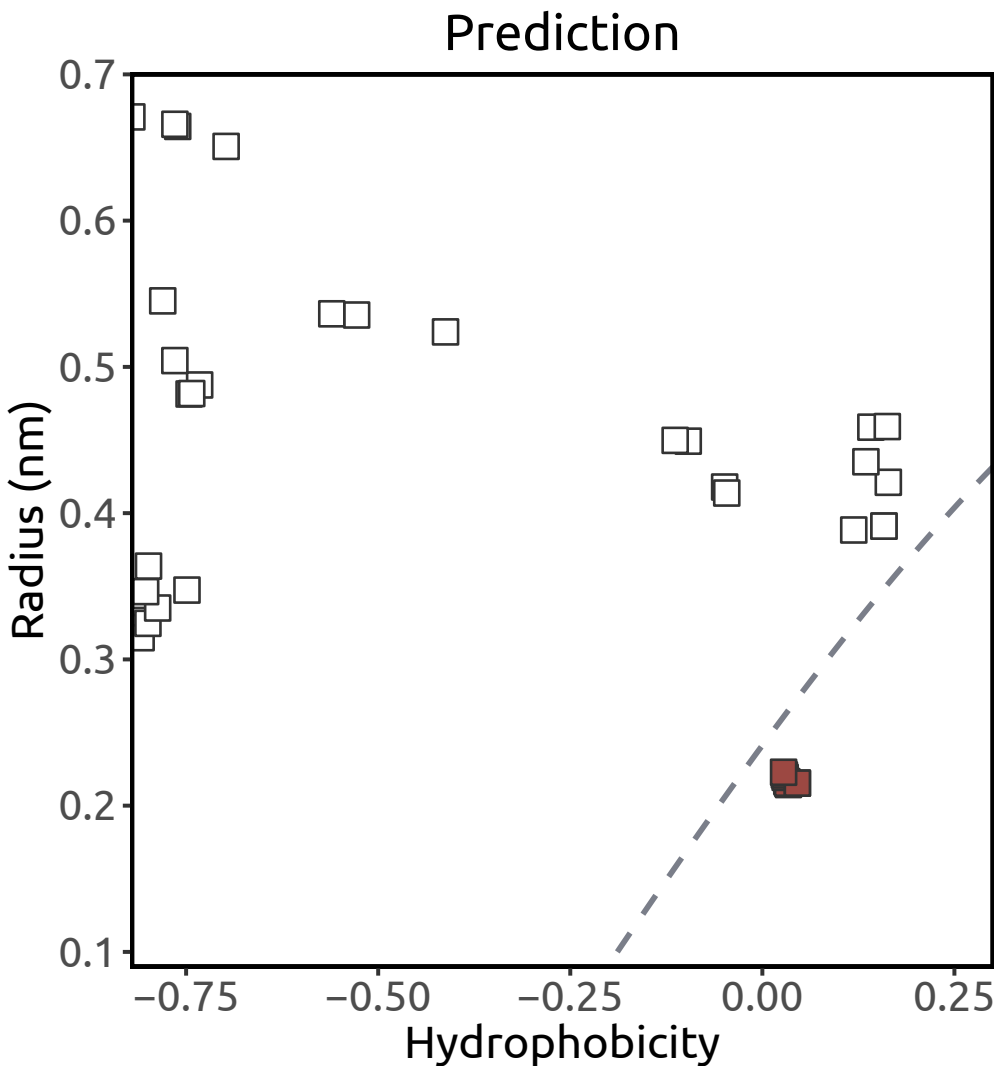
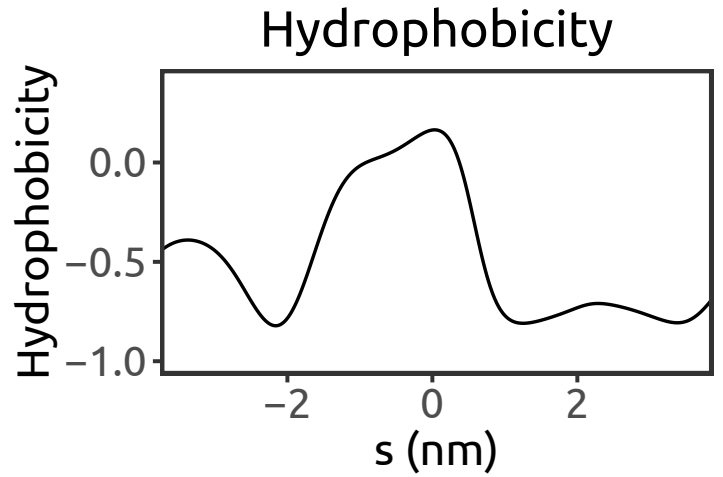
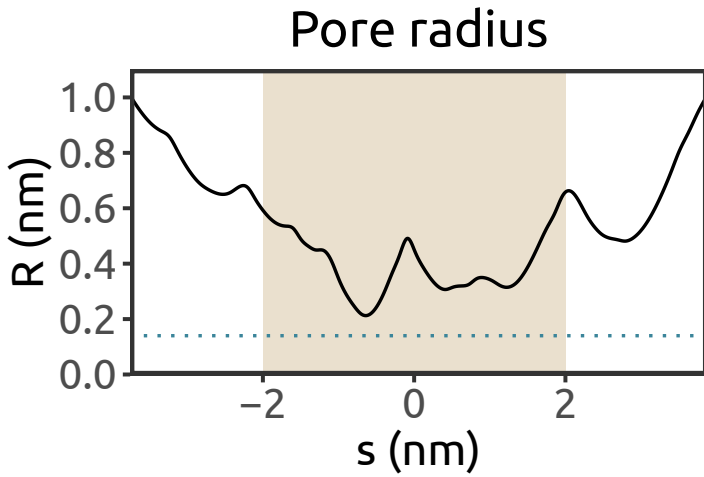
barrier to water

RyR2 (PDB ID: 5GOA)

Sus scrofa

cryo-EM (4.2 Å)

Peng et al., 2016



Heuristic score:
0.32 (n = 8)

Simulation result:
hydrated channel

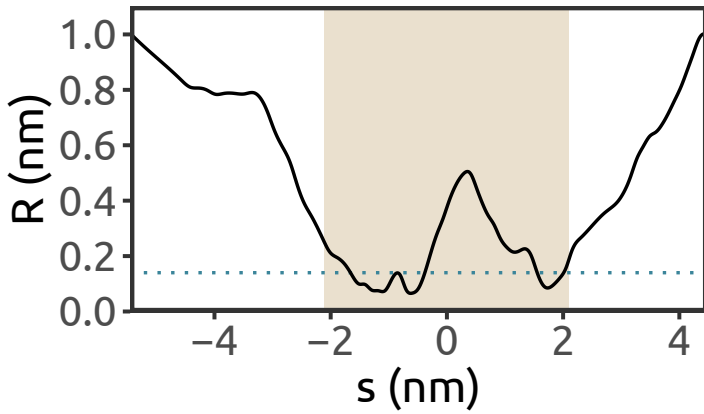
TPC1 (PDB ID: 5DQQ)

Arabidopsis thaliana

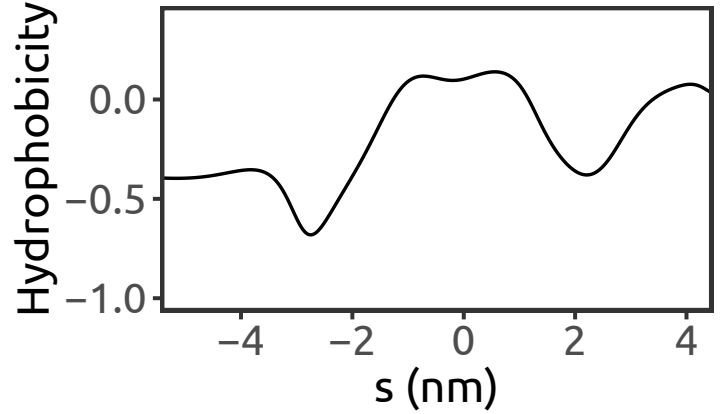
X-ray (2.87 Å)

Kintzer & Stroud, 2016

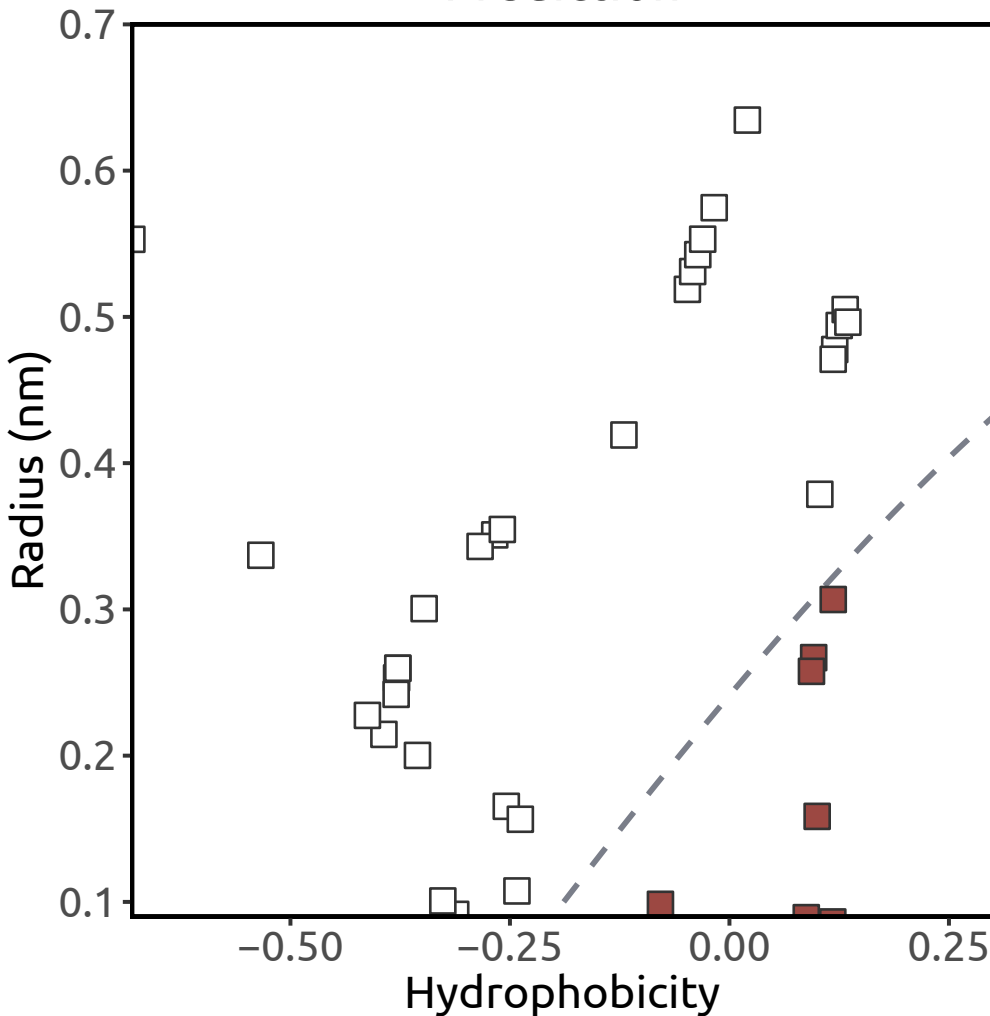
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.56 (n = 12)

Simulation result:
barrier to water

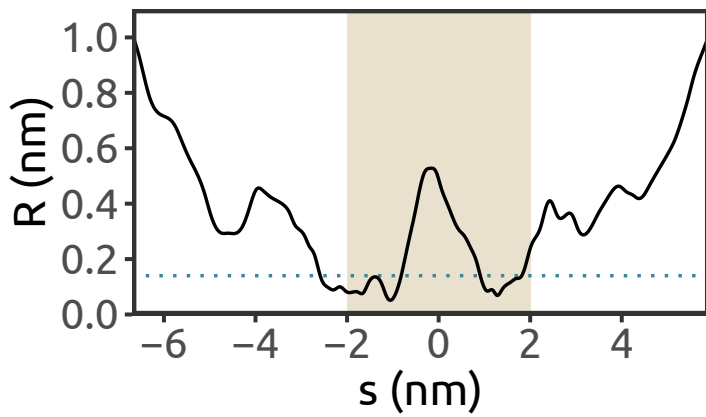
TPC1 (PDB ID: 6C96)

Mus musculus

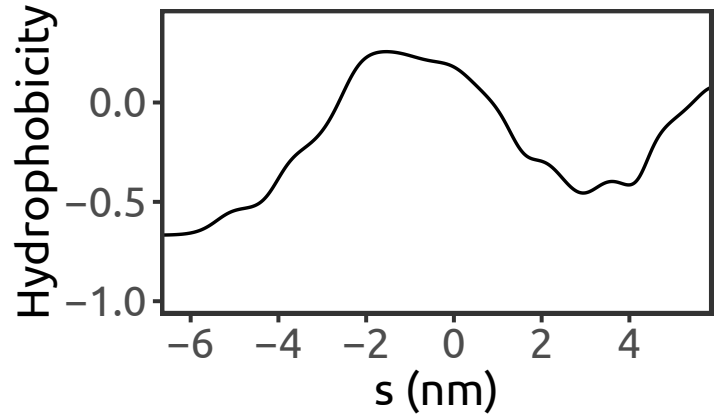
cryo-EM (3.4 Å)

She et al., 2018

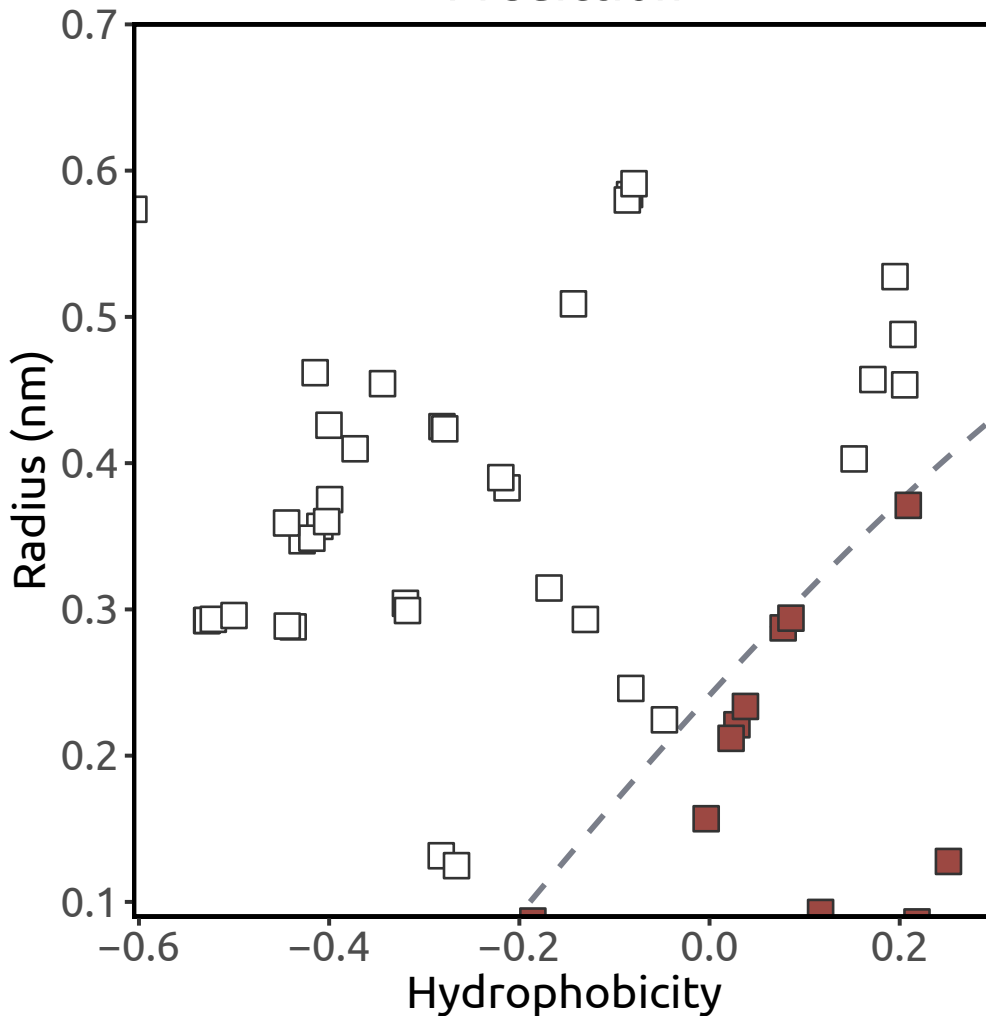
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.55 (n = 17)

Simulation result:
barrier to water

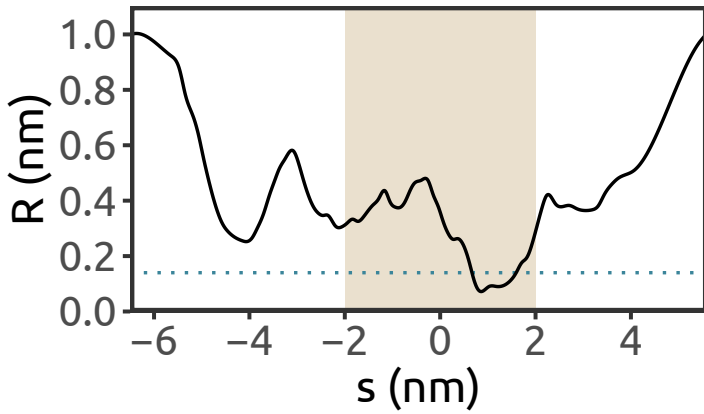
TPC1 (PDB ID: 6C9A)

Mus musculus

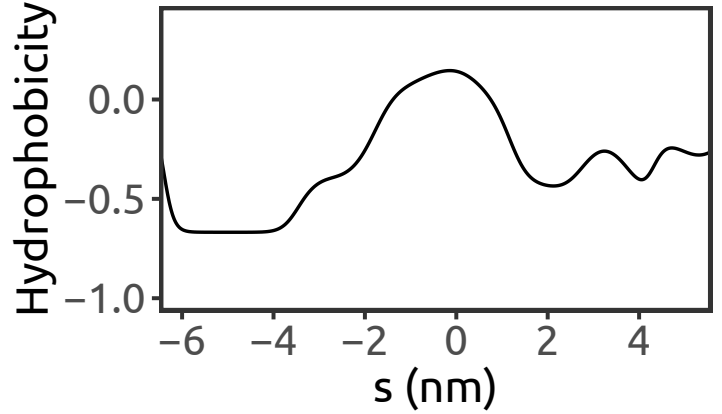
cryo-EM (3.2 Å)

She et al., 2018

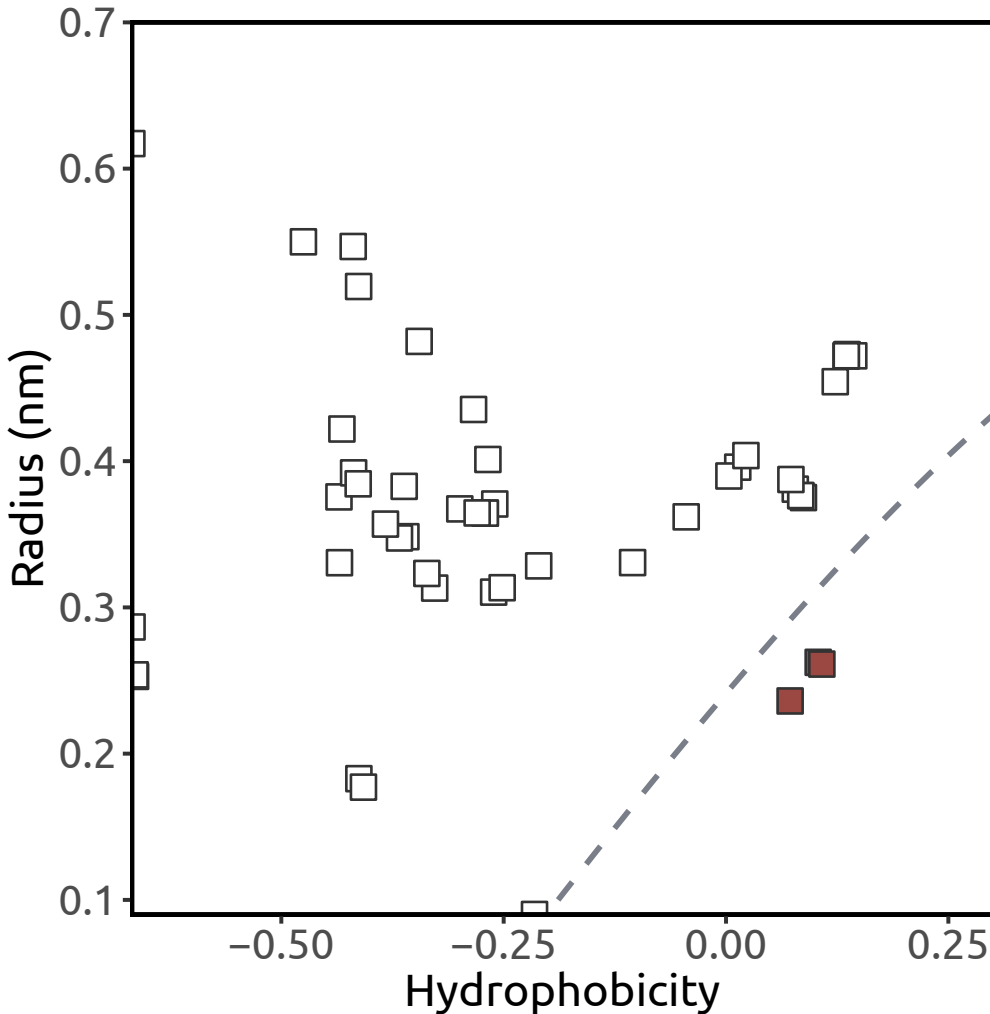
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.25 (n = 4)

Simulation result:
hydrated channel

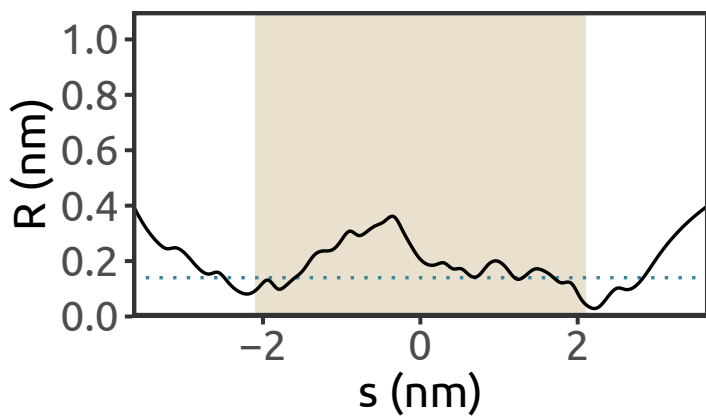
ASIC1 (PDB ID: 2QTS)

Gallus gallus

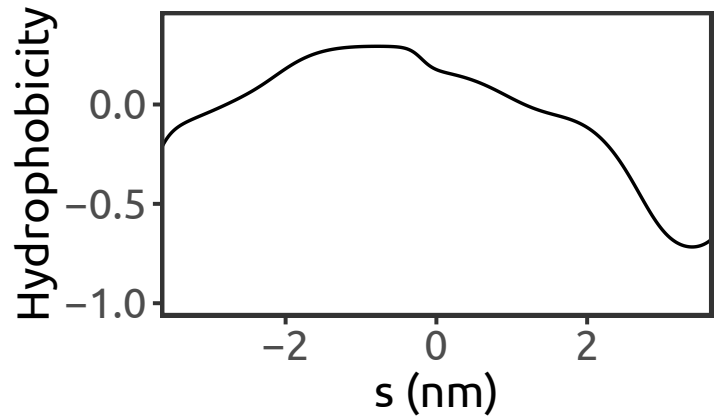
X-ray (1.9 Å)

Jasti et al., 2007

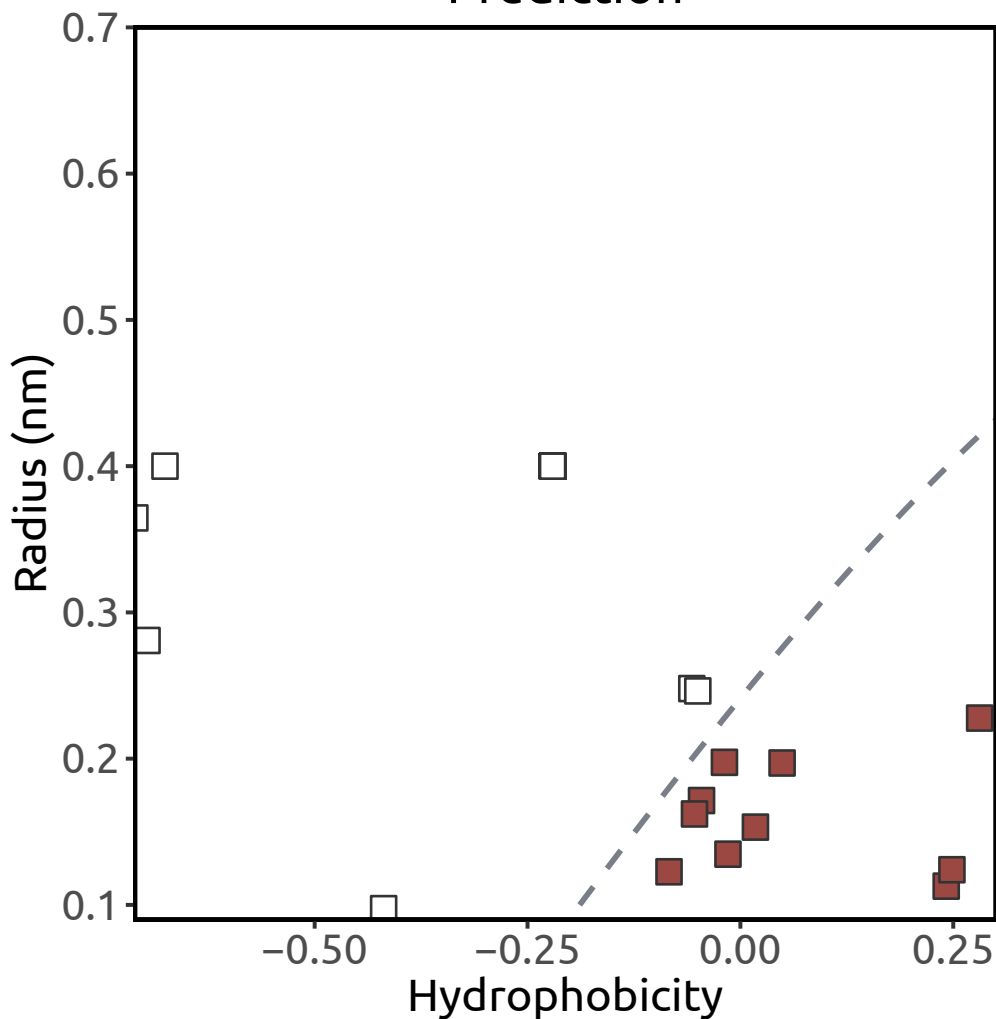
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.35 (n = 13)

Simulation result:
barrier to water

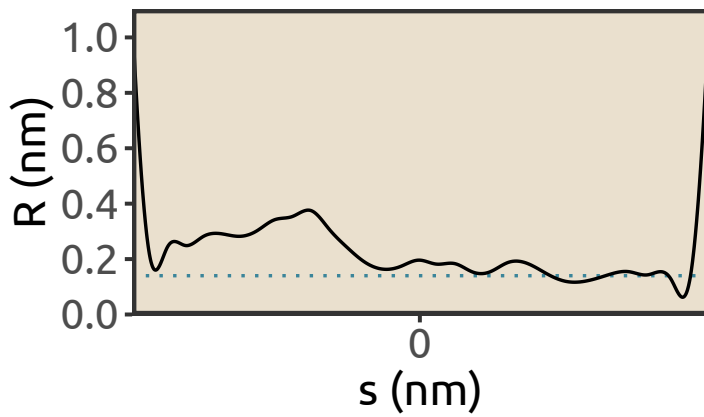
ASIC1 (PDB ID: 3S3W)

Gallus gallus

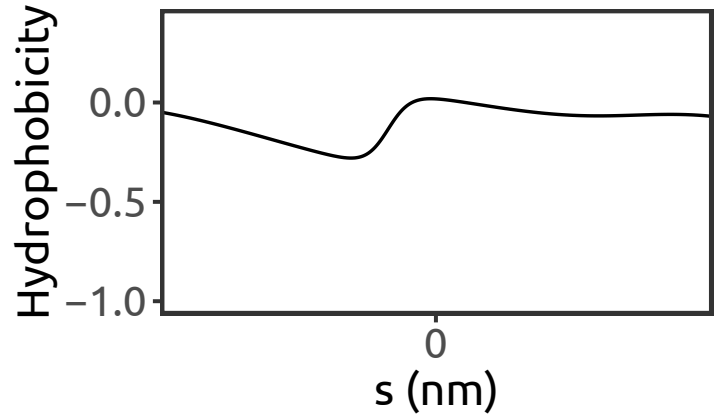
X-ray (2.6 Å)

Dawson et al., 2012

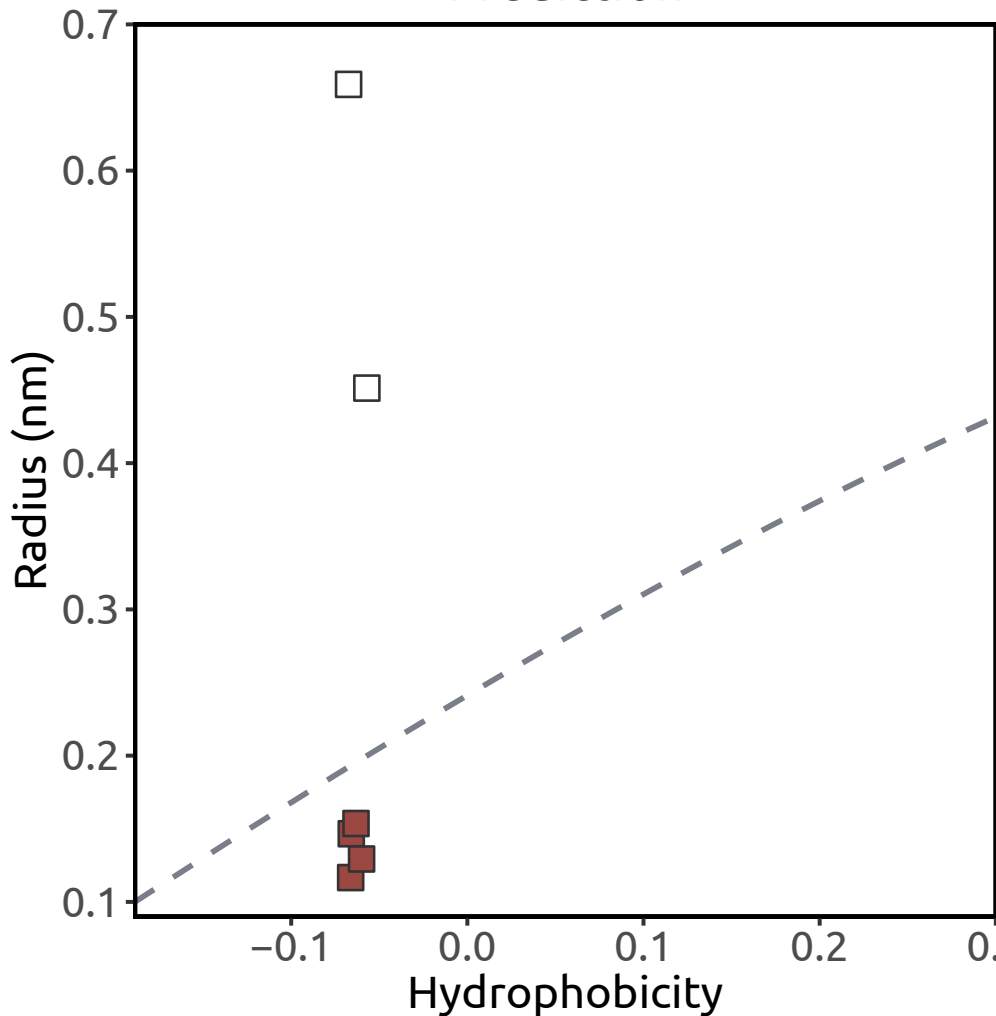
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.19 (n = 4)

Simulation result:
barrier to water

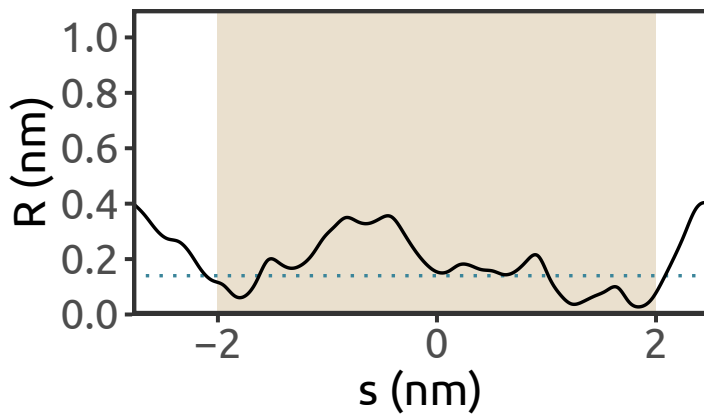
ASIC1 (PDB ID: 4NTW)

Gallus gallus

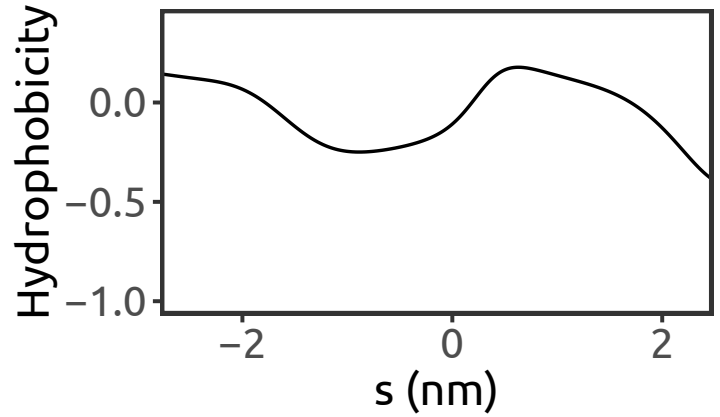
X-ray (2.07 Å)

Baconguis et al., 2014

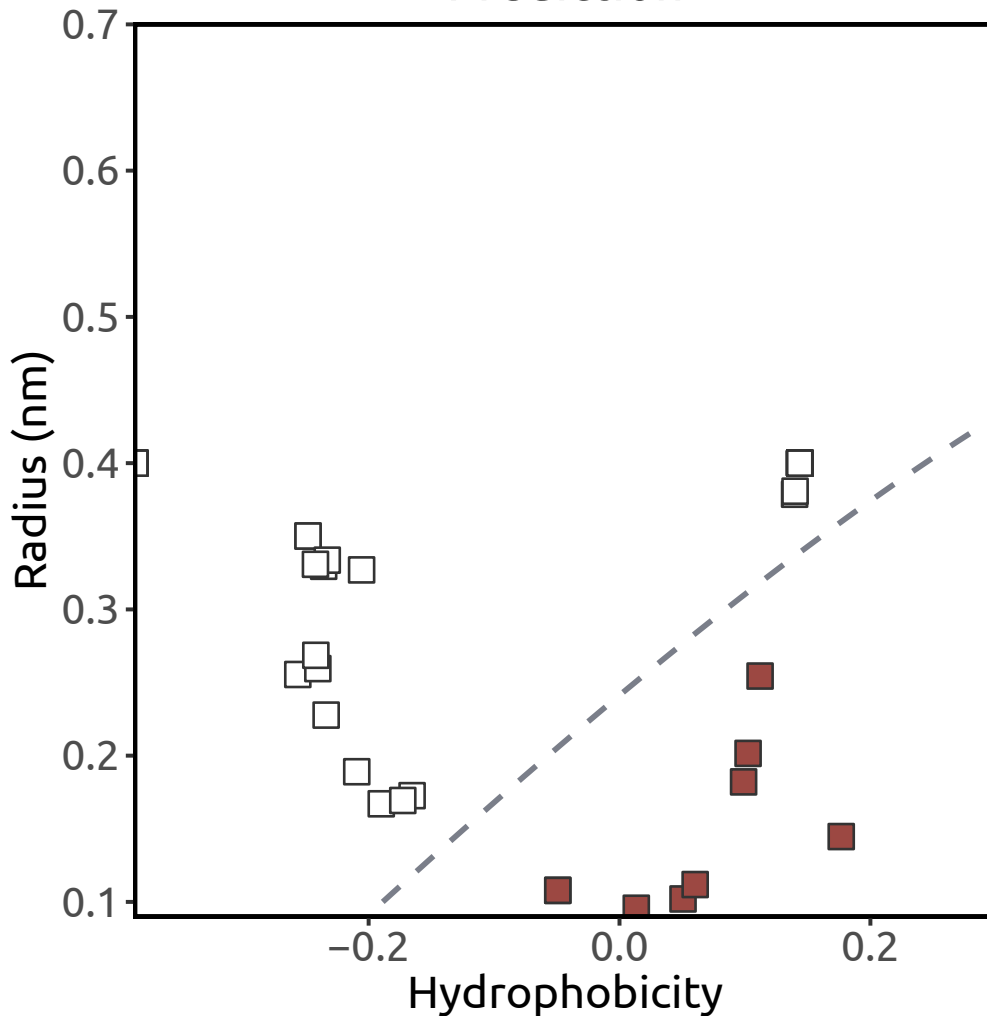
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.35 (n = 17)

Simulation result:
hydrated channel

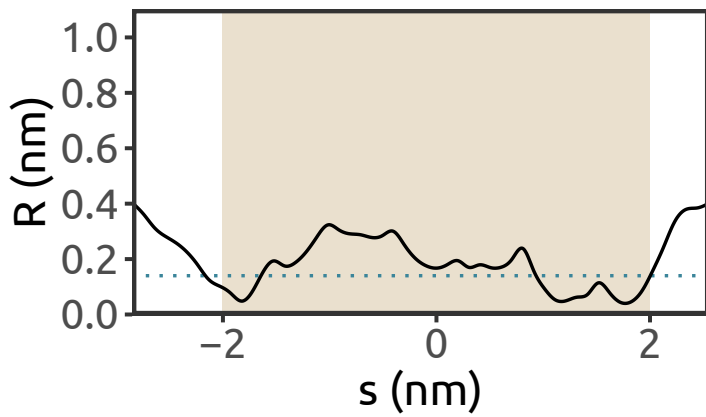
ASIC1 (PDB ID: 4NTY)

Gallus gallus

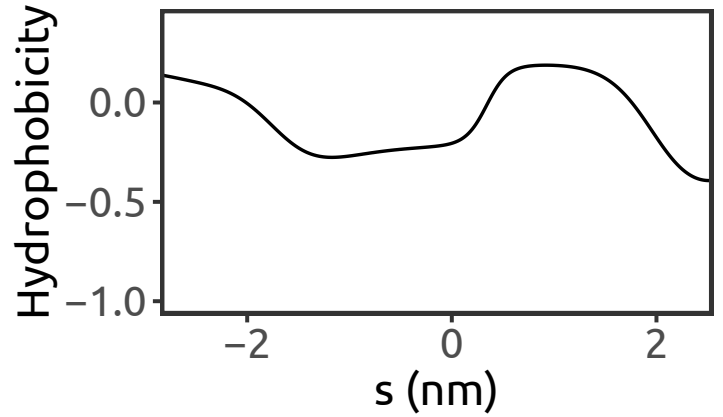
X-ray (2.65 Å)

Baconguis et al., 2014

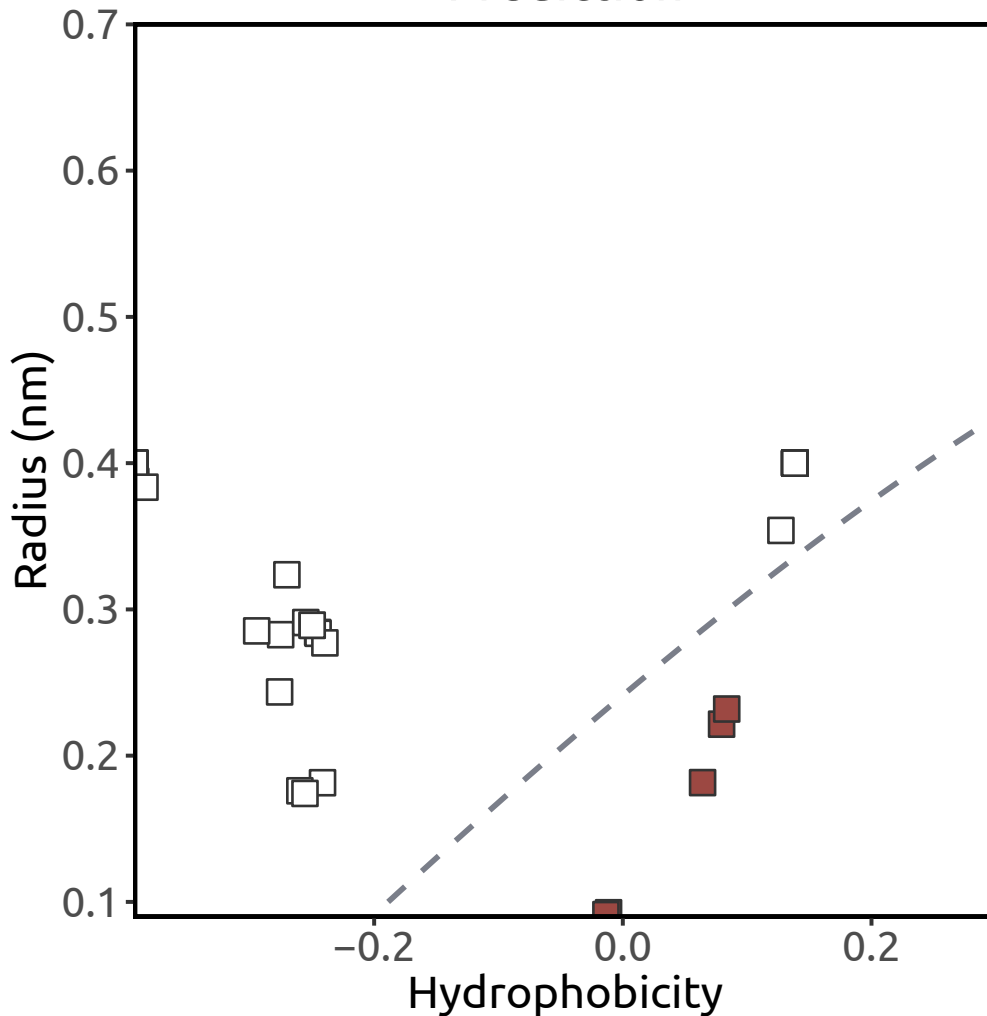
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.83 (n = 12)

Simulation result:

hydrated channel

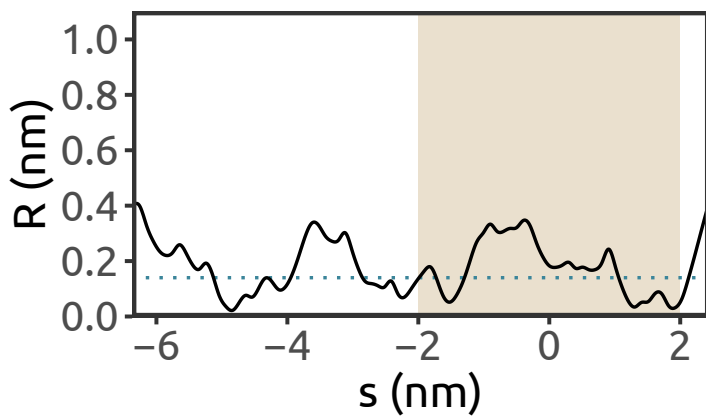
ASIC1 (PDB ID: 4NYK)

Gallus gallus

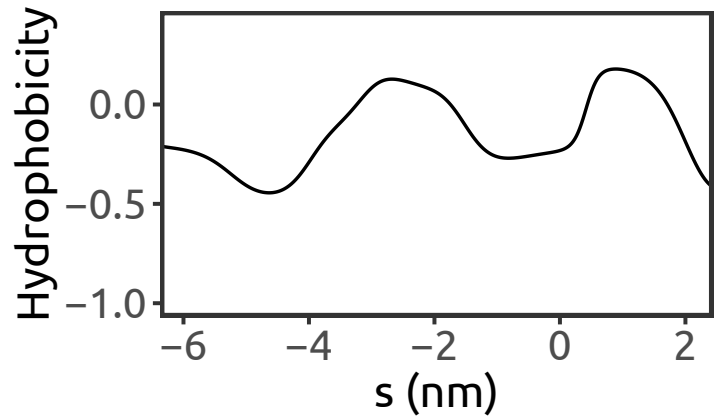
X-ray (3 Å)

Gonzales et al., 2009

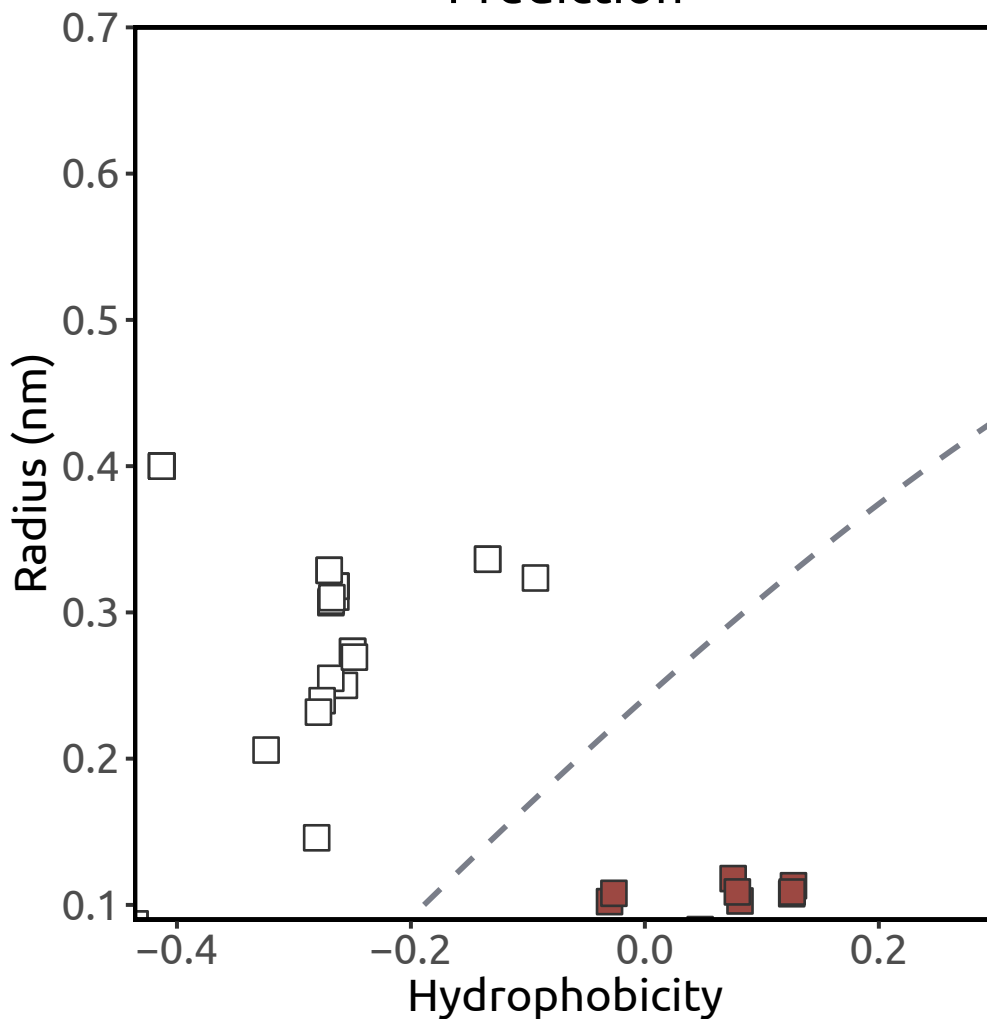
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.43 (n = 15)

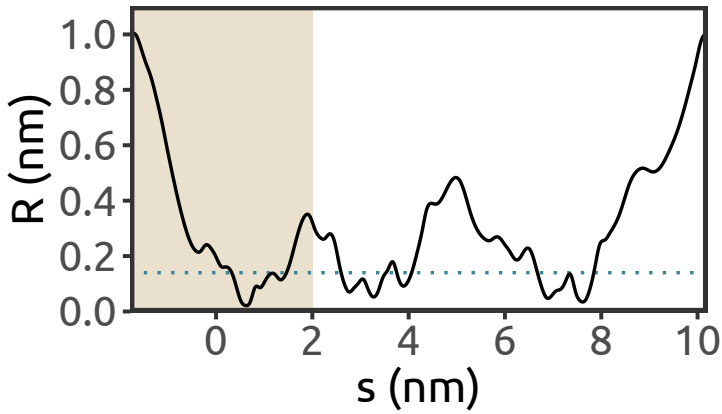
Simulation result:
barrier to water

ASIC1 (PDB ID: 5WKU)

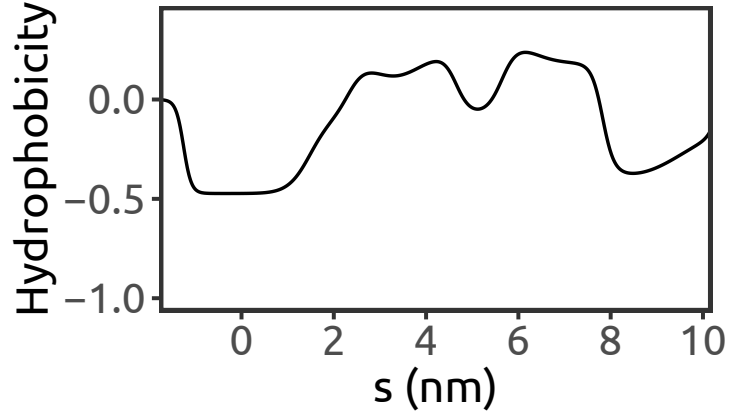
Gallus gallus
X-ray (2.95 Å)

Yoder et al., 2018

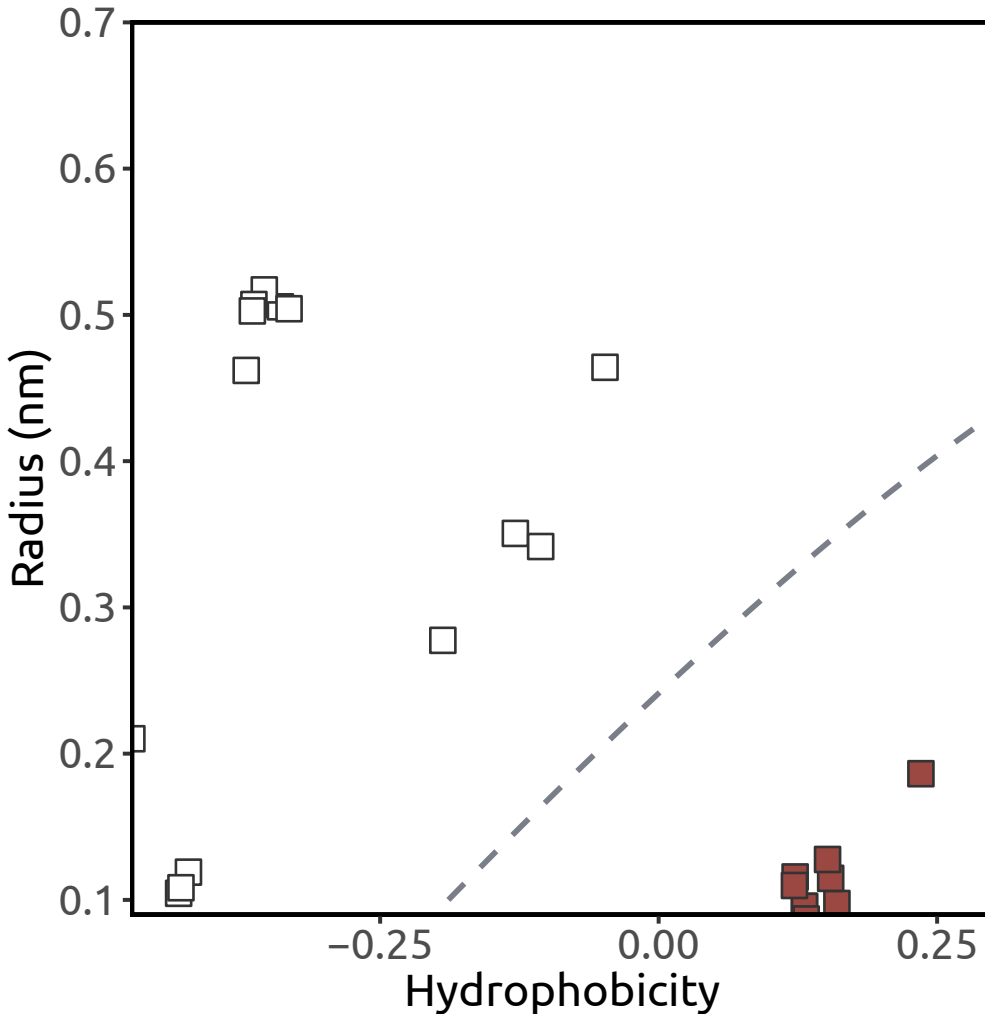
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.67 (n = 13)

Simulation result:
barrier to water

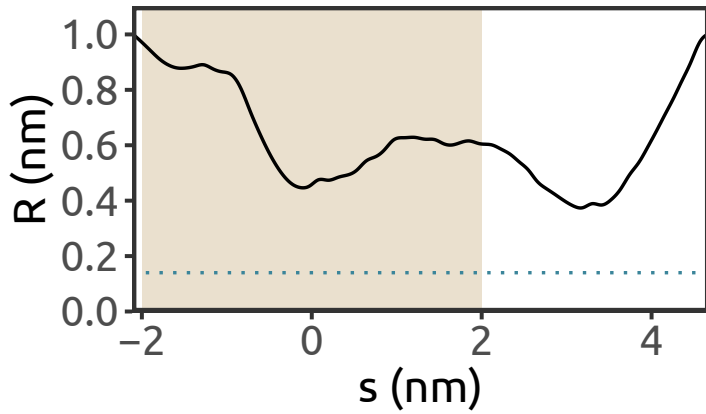
ExbBExbD (PDB ID: 5SV0)

Escherichia coli

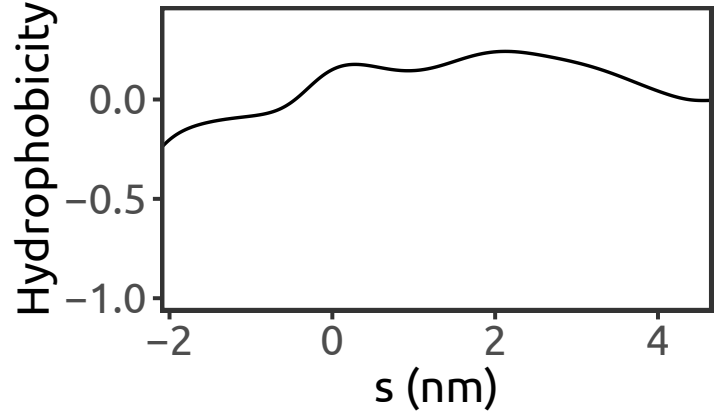
X-ray (2.6 Å)

Celia et al., 2016

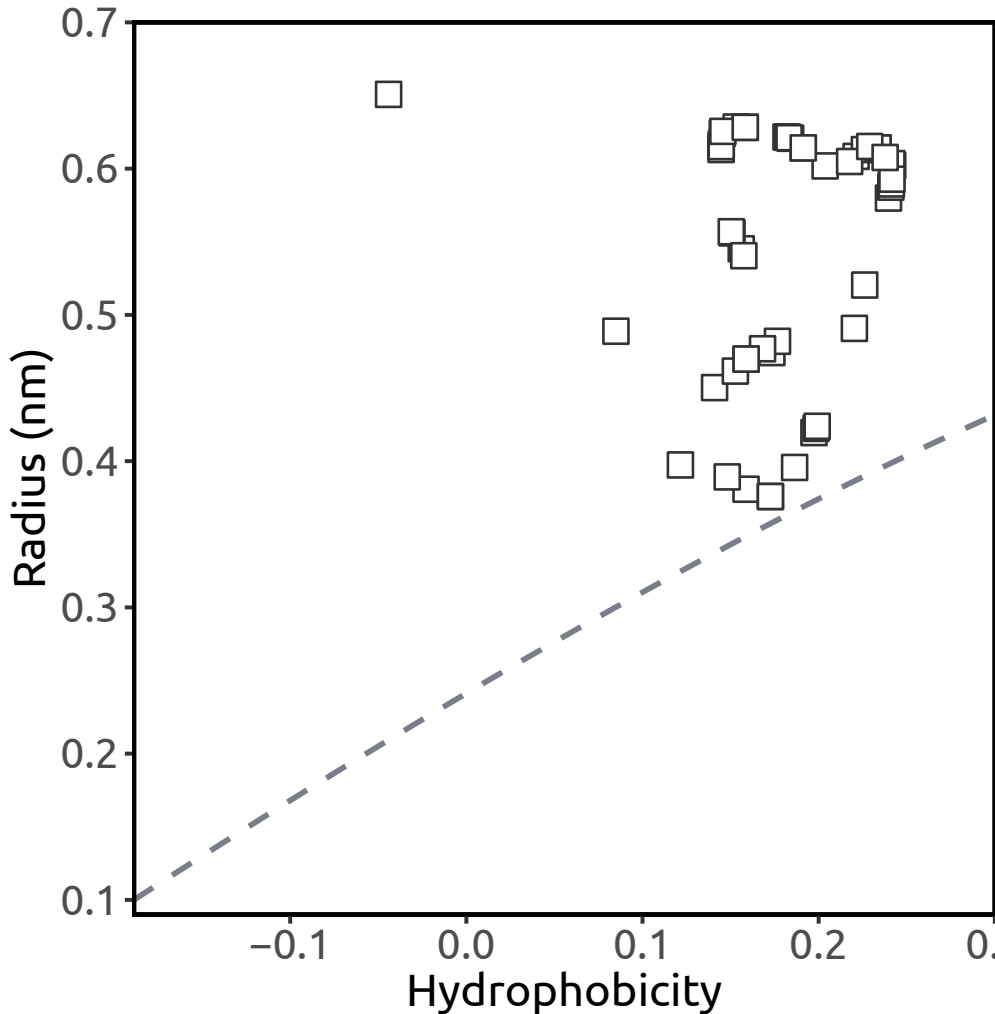
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

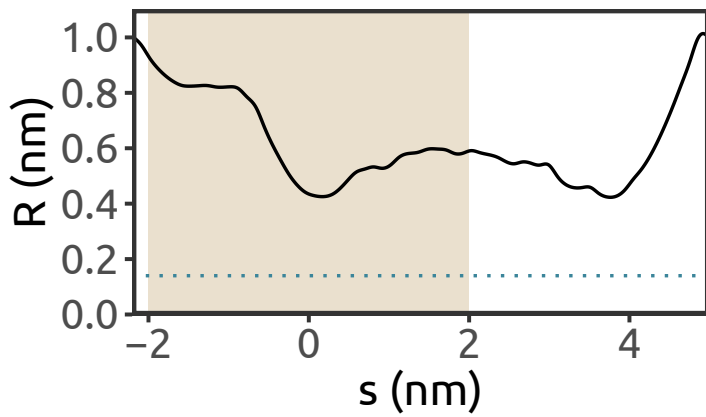
ExbBExbD (PDB ID: 5SV1)

Escherichia coli

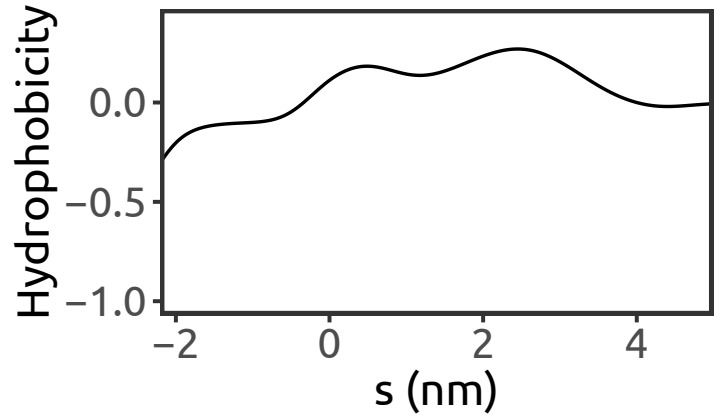
X-ray (3.5 Å)

Celia et al., 2016

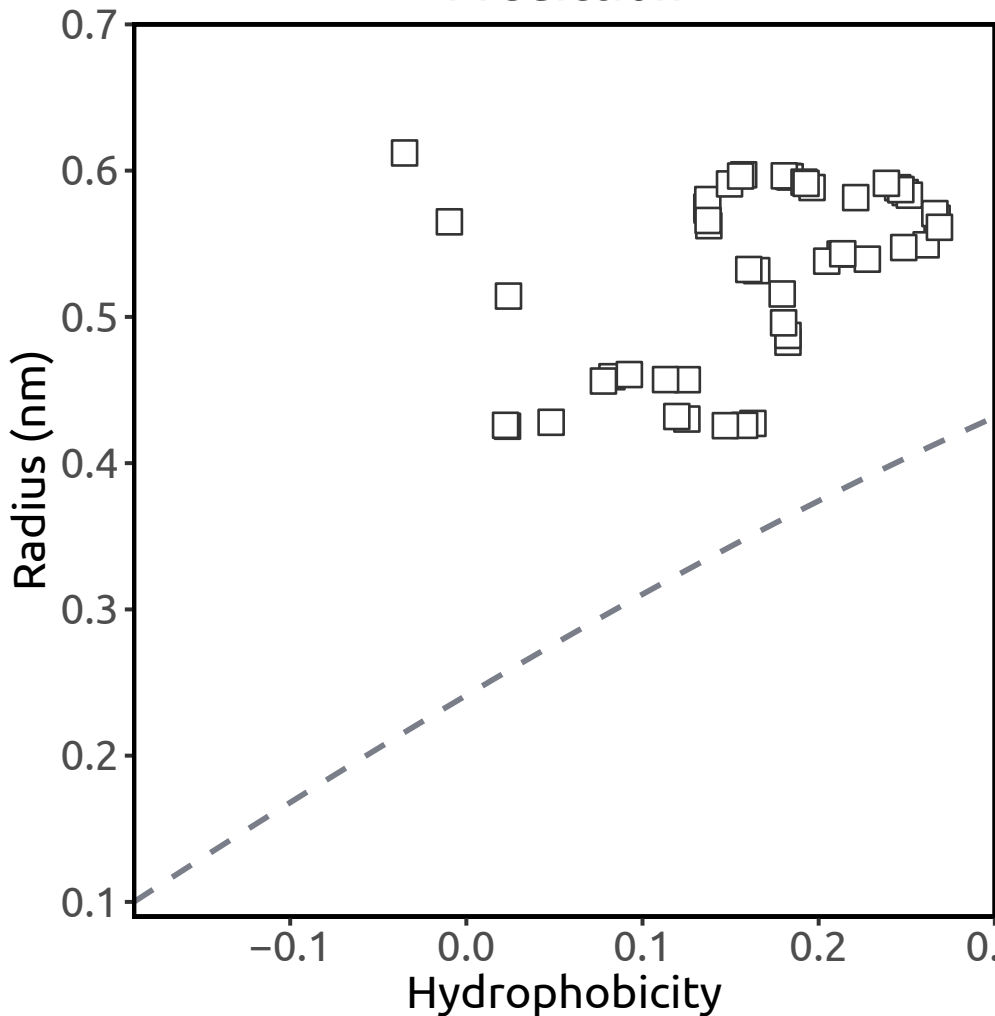
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

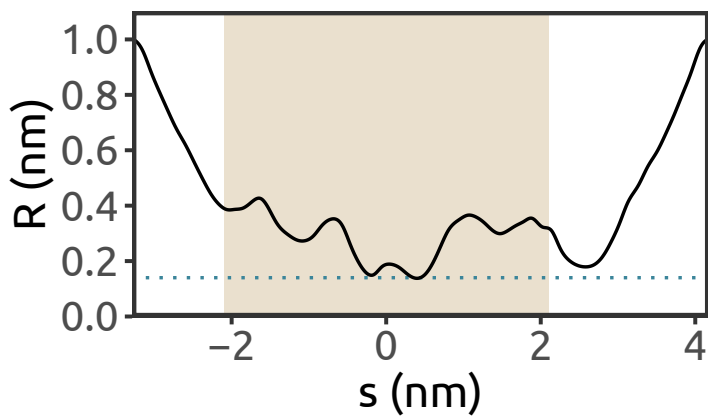
M2 (PDB ID: 2KIX)

Influenza B

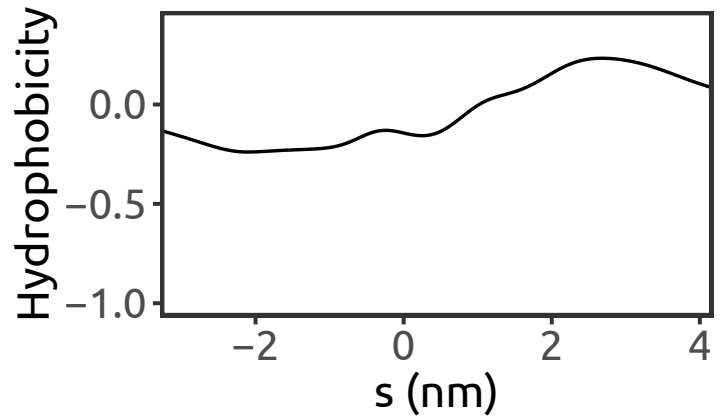
aq NMR

Wang et al., 2009

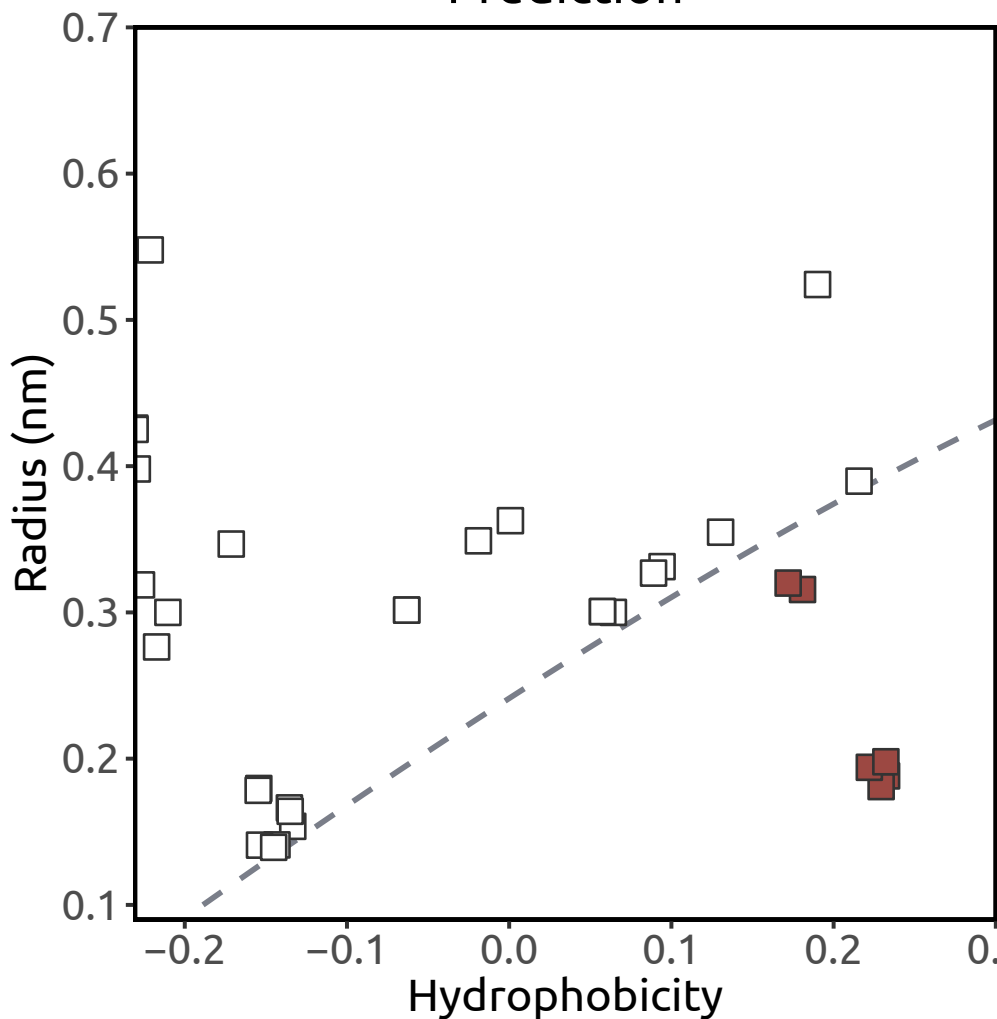
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.76 (n = 6)

Simulation result:
barrier to water

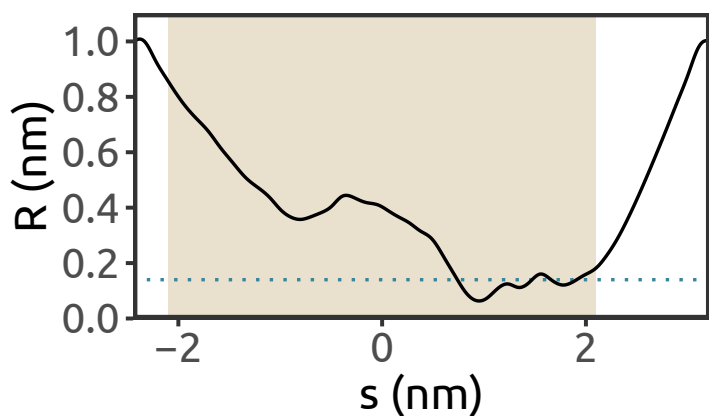
M2 (PDB ID: 3BKD)

Influenza A

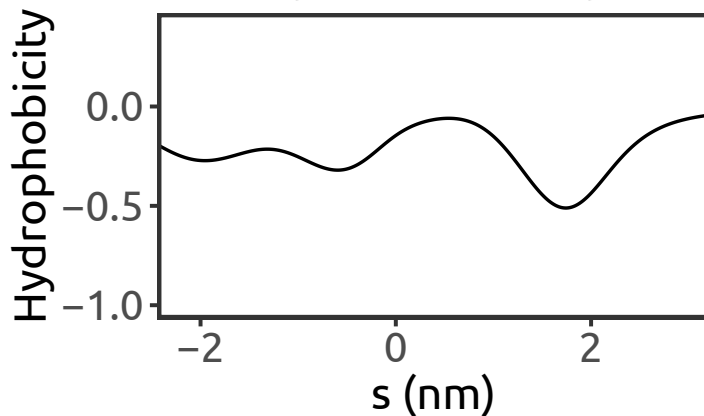
X-ray (2.05 Å)

Stouffer et al., 2008

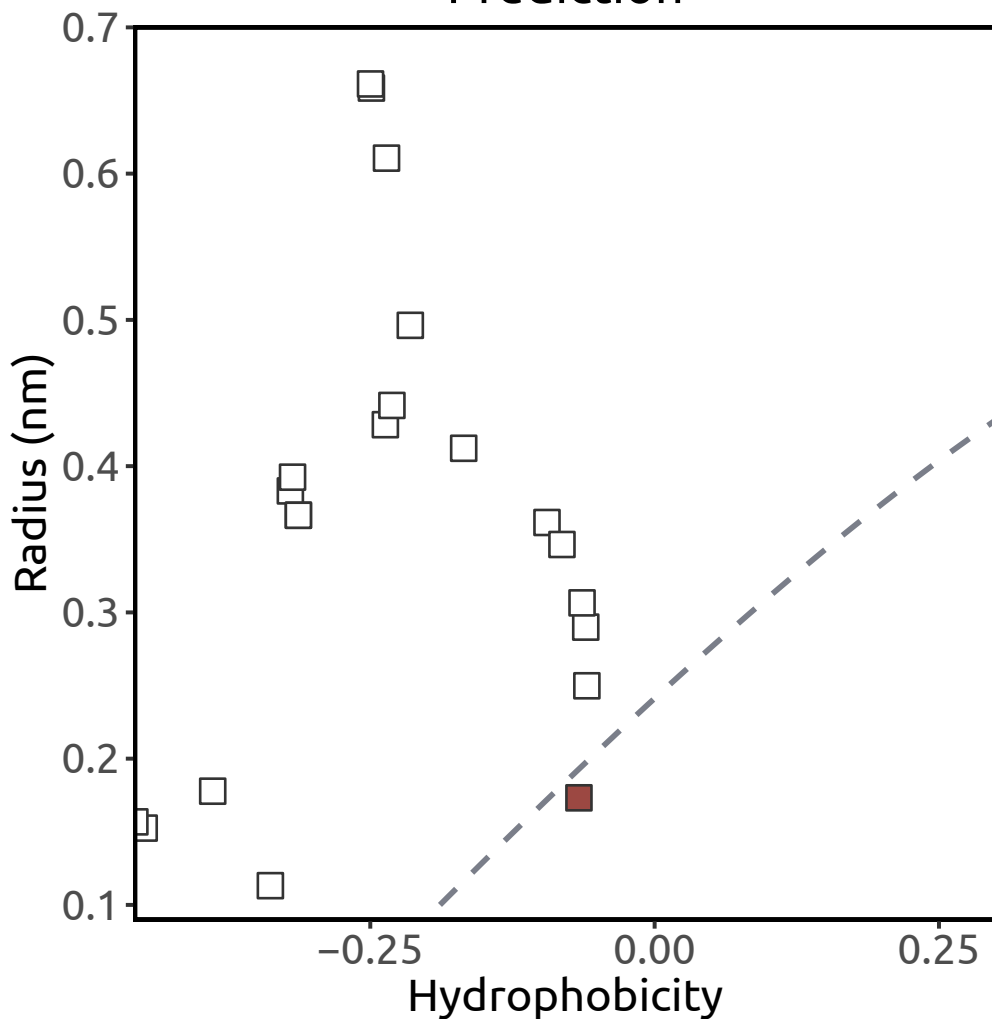
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.27 (n = 5)

Simulation result:
barrier to water

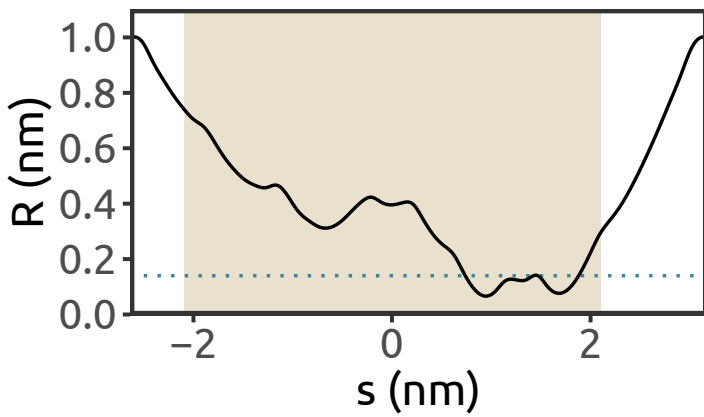
M2 (PDB ID: 4QKC)

Influenza A

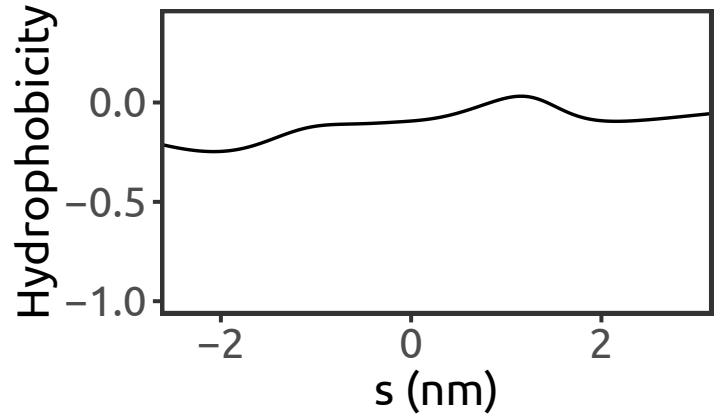
X-ray (1.1 Å)

Thomaston et al., 2015

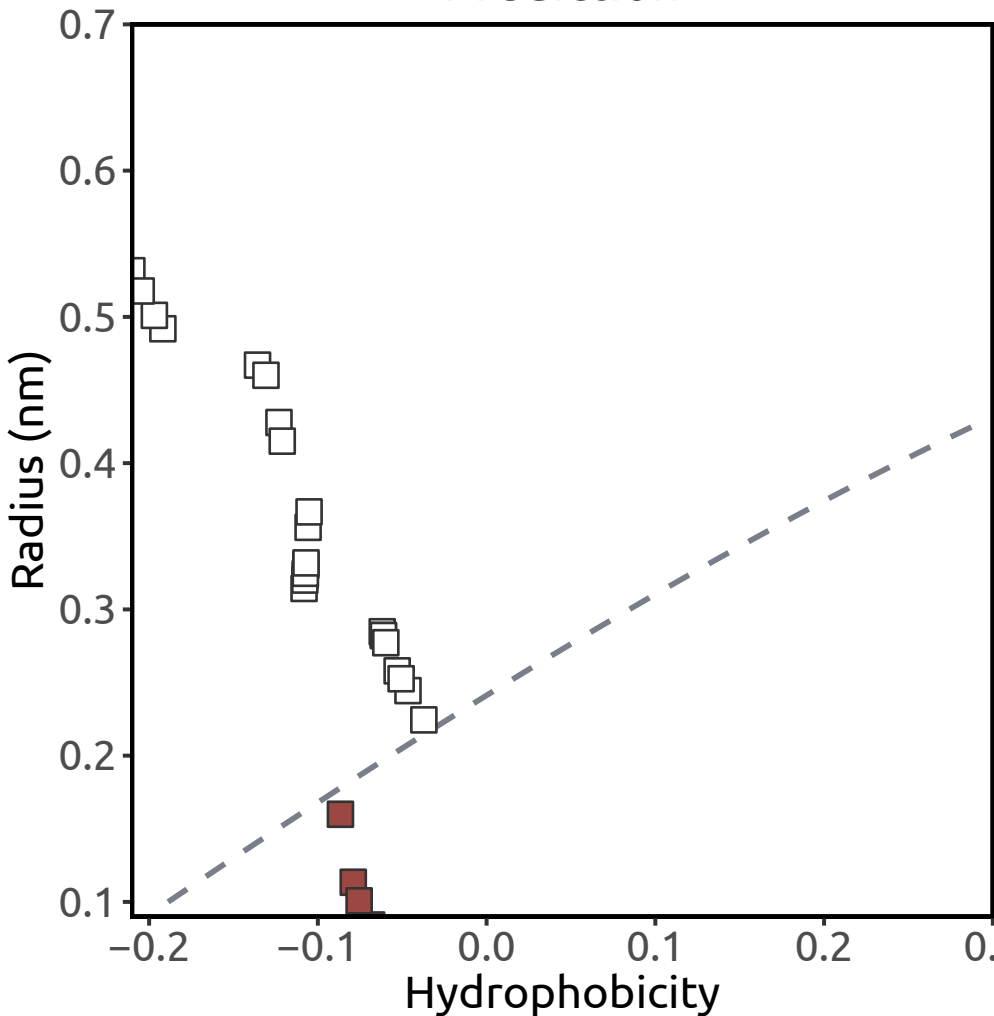
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.82 (n = 8)

Simulation result:

barrier to water

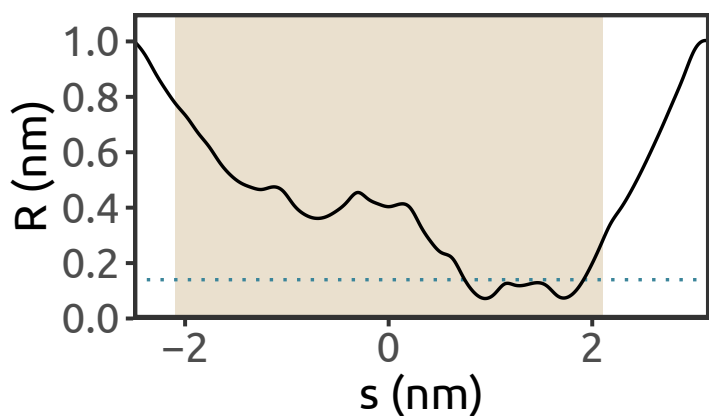
M2 (PDB ID: 5J00)

Influenza A

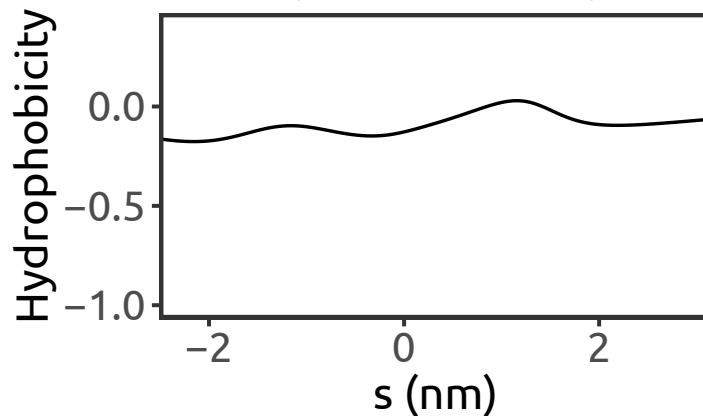
X-ray (1.41 Å)

Thomaston et al., 2017

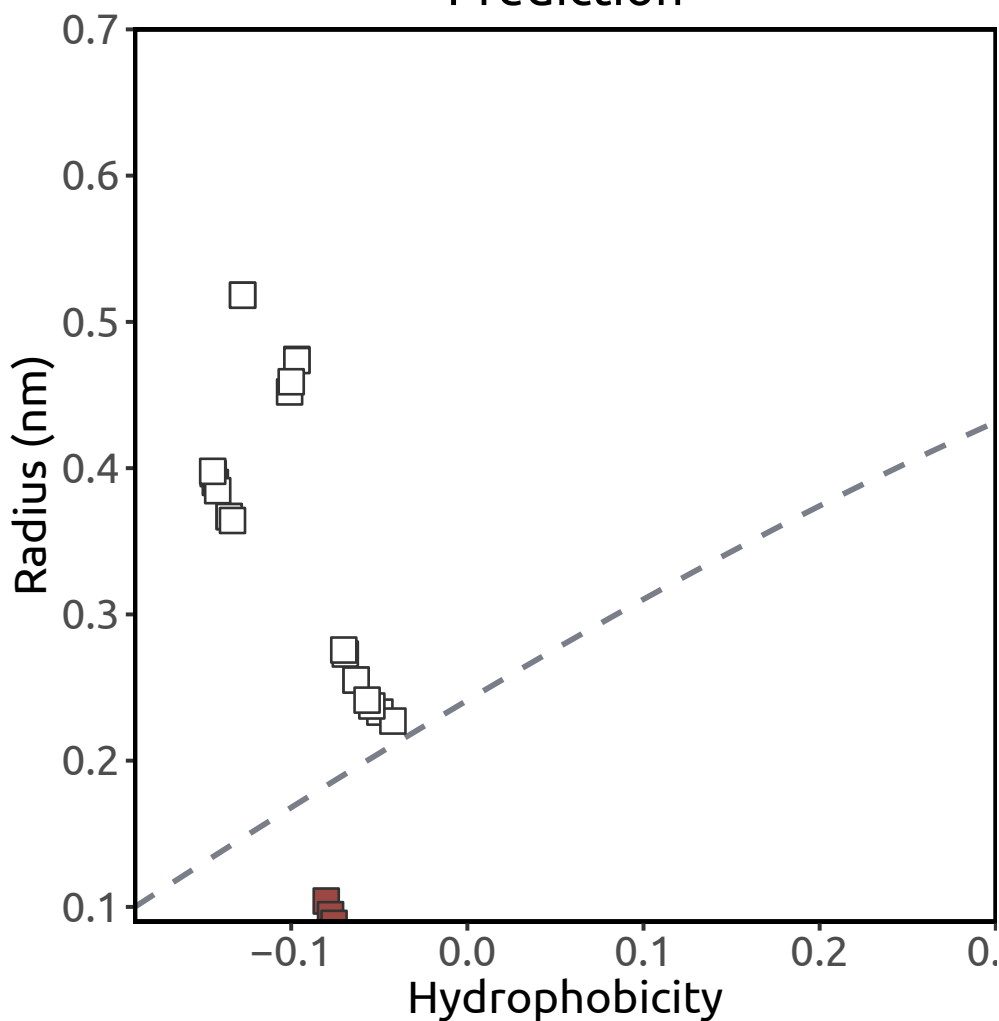
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.88 (n = 8)

Simulation result:

barrier to water

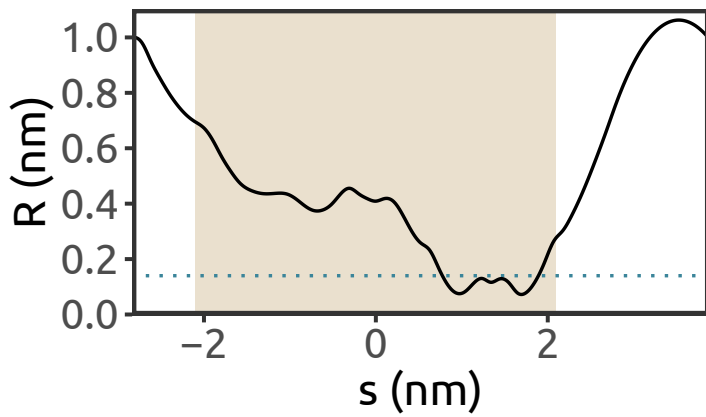
M2 (PDB ID: 5TTC)

Influenza A

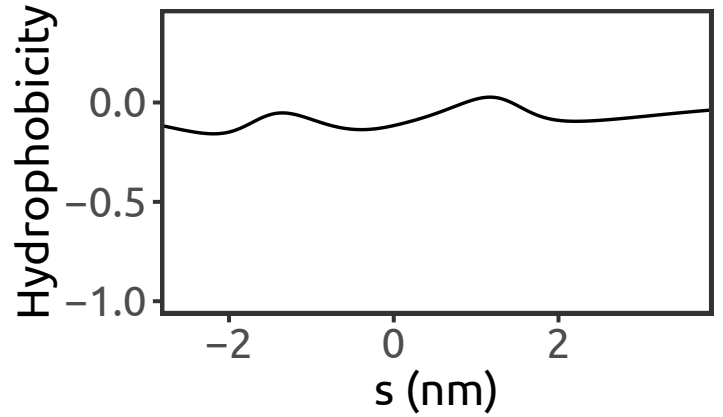
X-ray (1.4 Å)

Thomaston et al., 2017

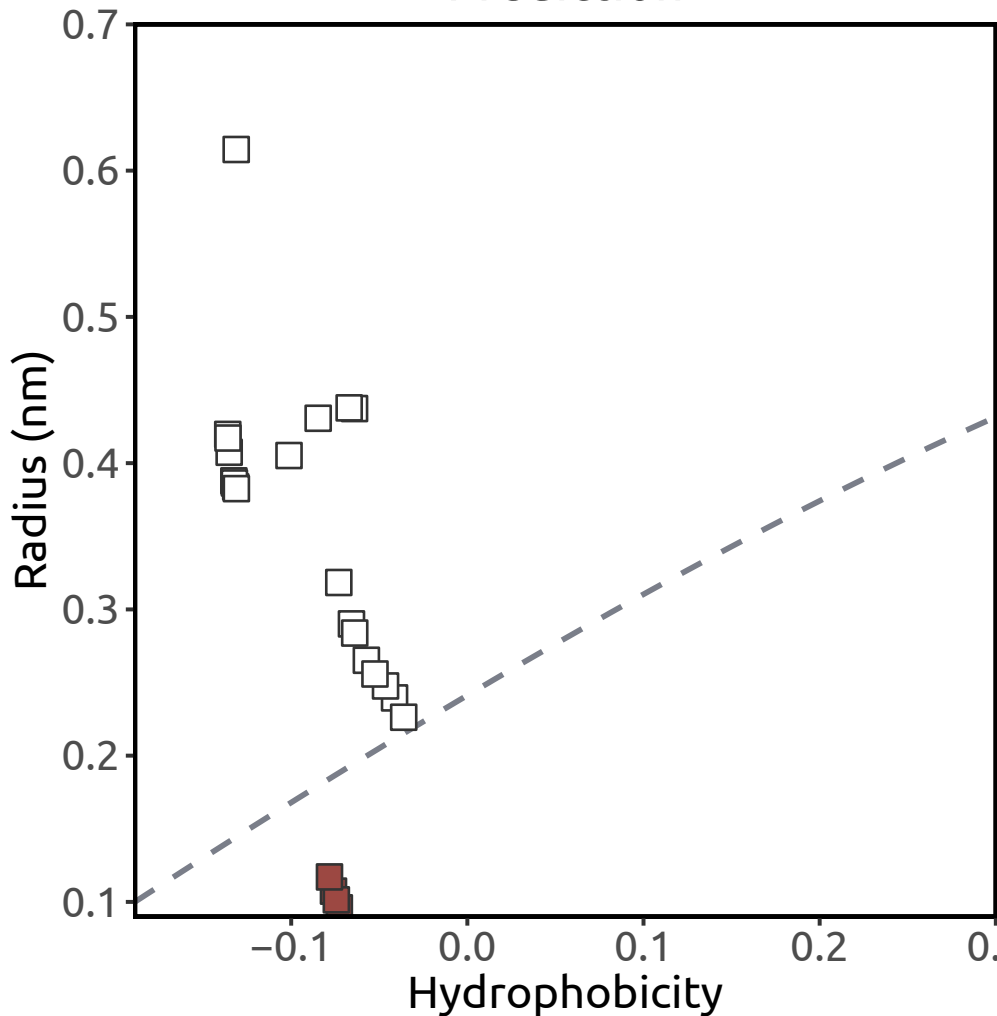
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.83 (n = 8)

Simulation result:

barrier to water

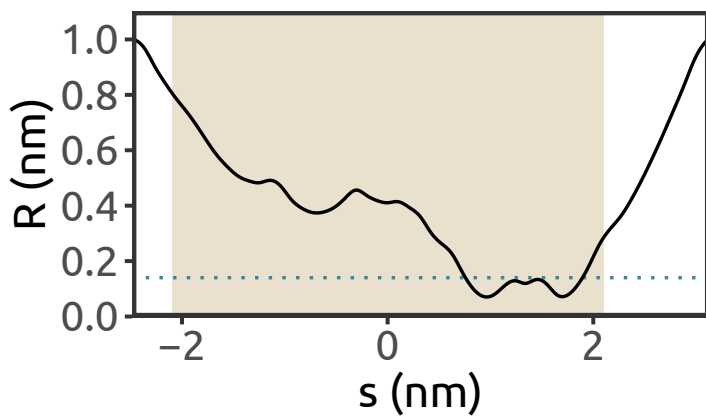
M2 (PDB ID: 5UM1)

Influenza A

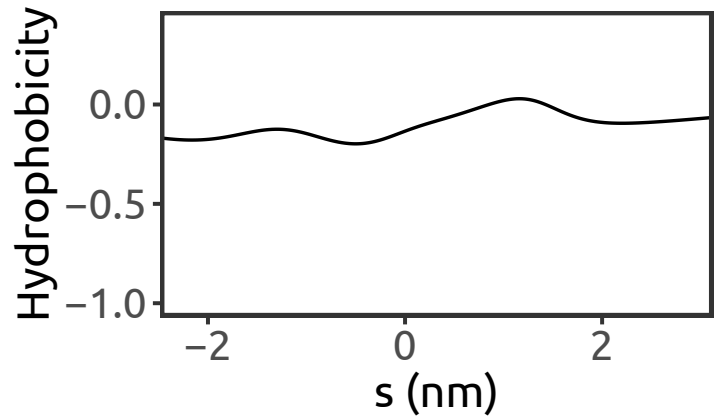
X-ray (1.45 Å)

Thomaston et al., 2017

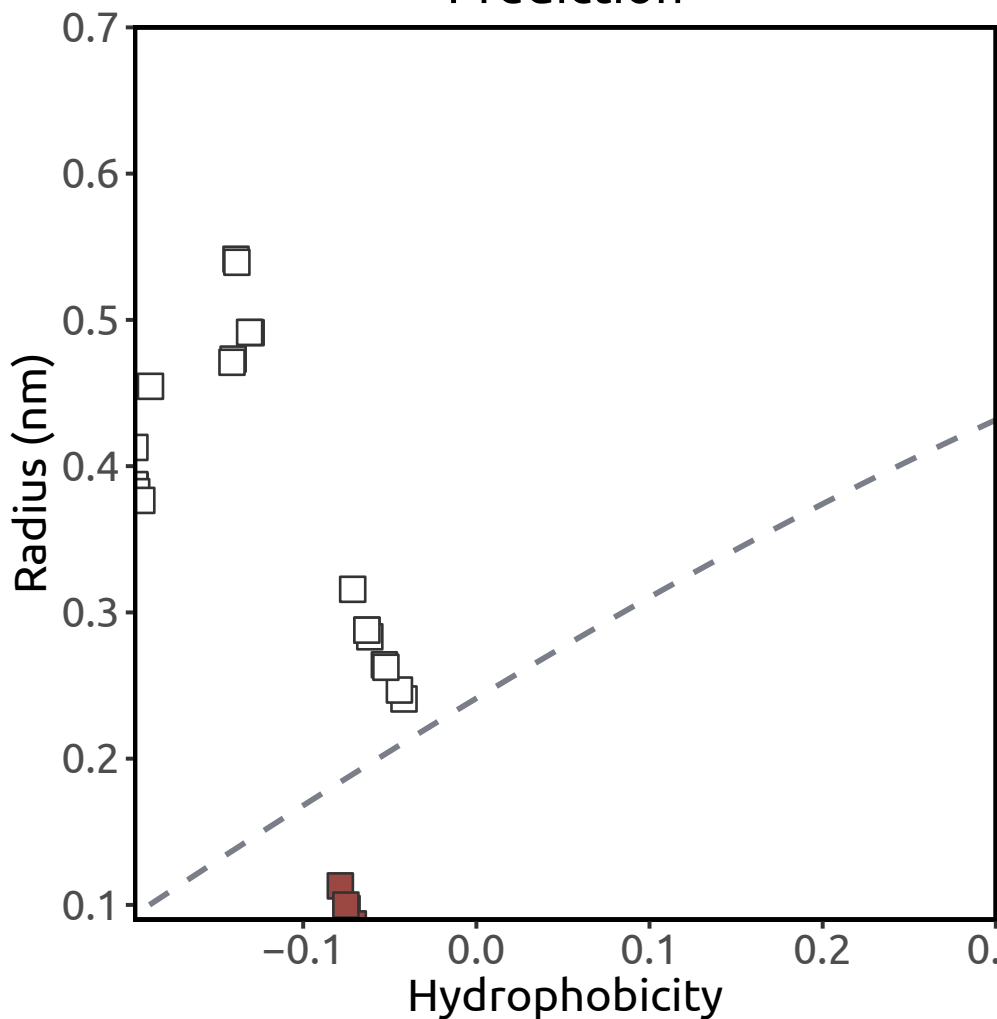
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.87 (n = 8)

Simulation result:
barrier to water

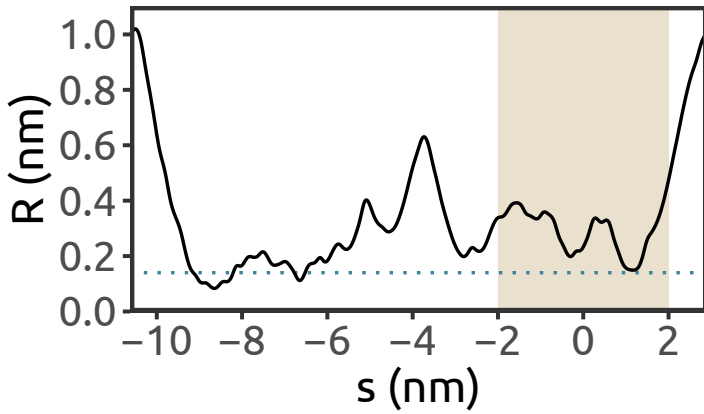
MCU (PDB ID: 5ID3)

Caenorhabditis elegans

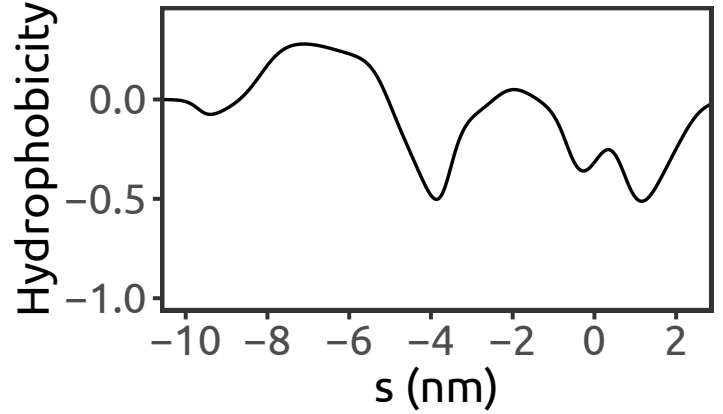
aq NMR

Oxenoid et al., 2016

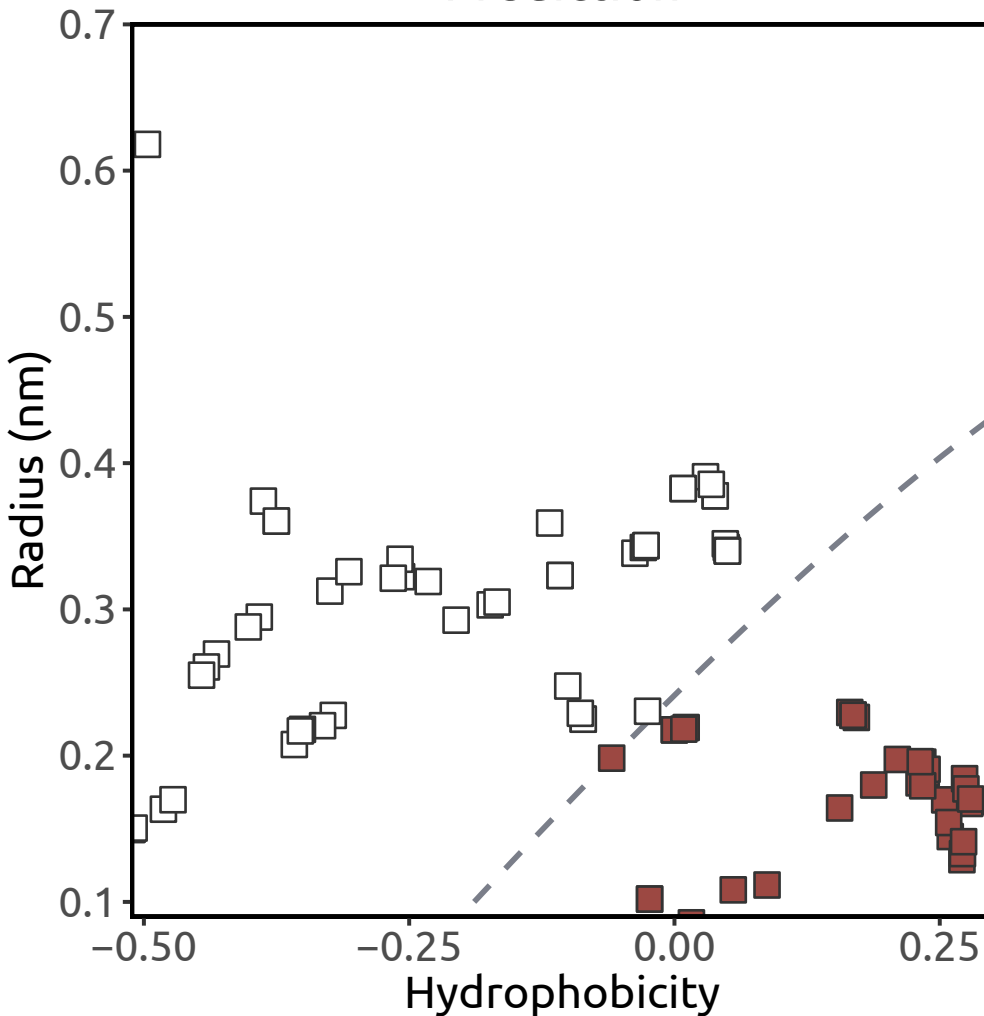
Pore radius



Hydrophobicity



Prediction



Heuristic score:

5.2 (n = 35)

Simulation result:

barrier to water

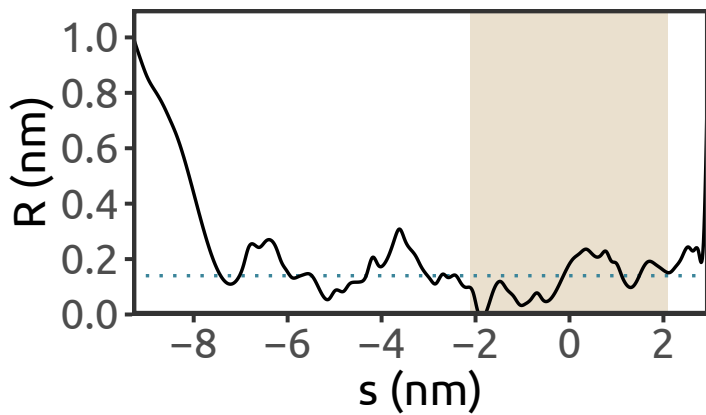
MgtE (PDB ID: 2ZY9)

Thermus thermophilus

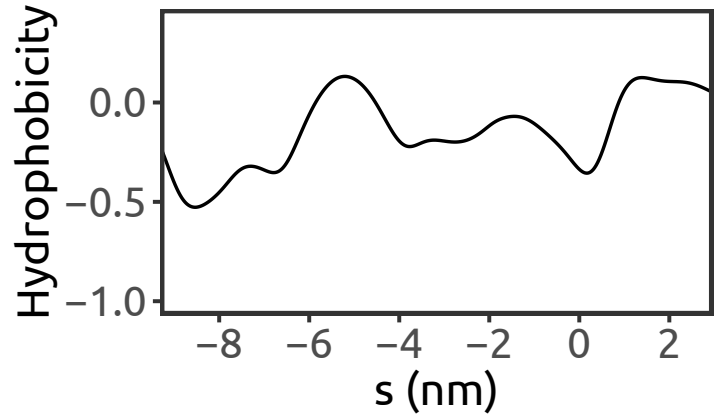
X-ray (2.94 Å)

Hattori et al., 2009

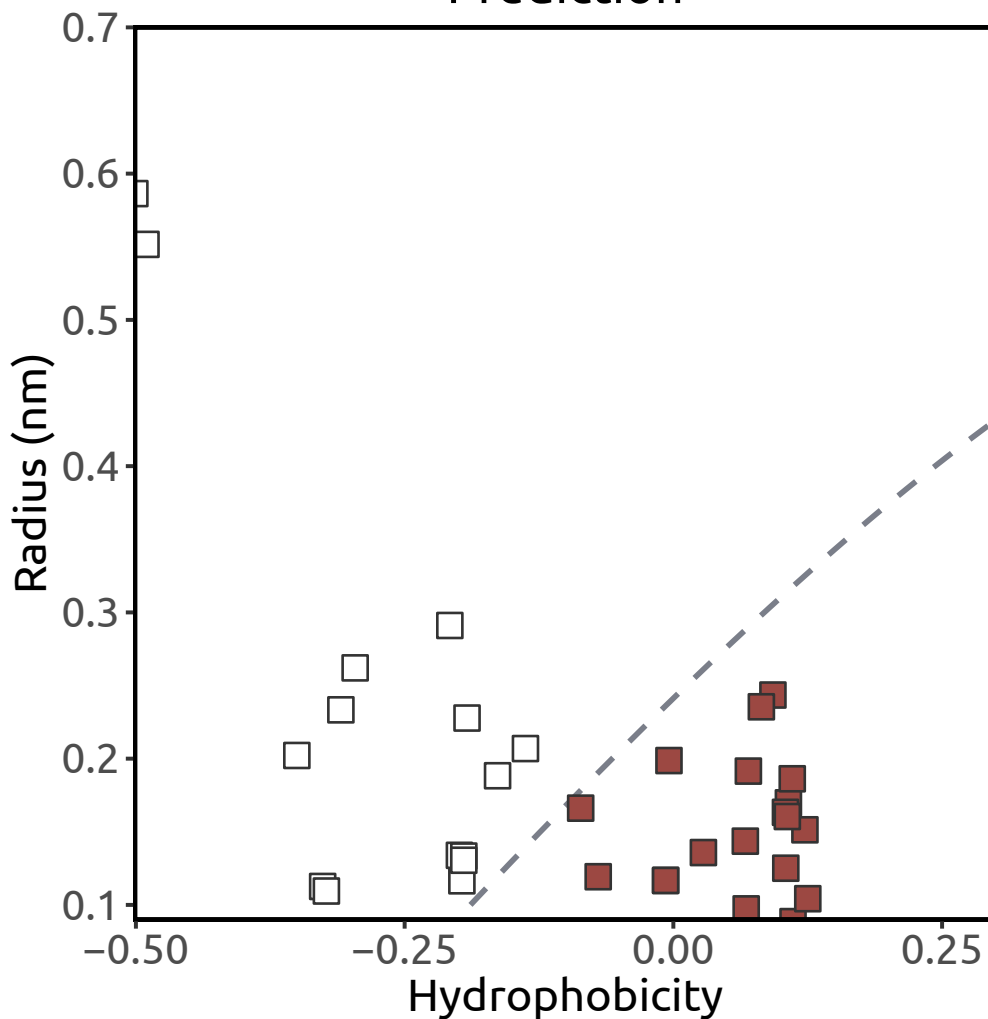
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.58 (n = 23)

Simulation result:
barrier to water

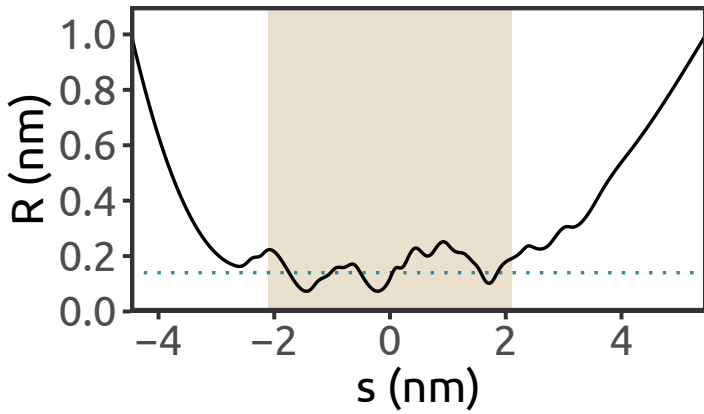
MgtE (PDB ID: 4U9N)

Thermus thermophilus

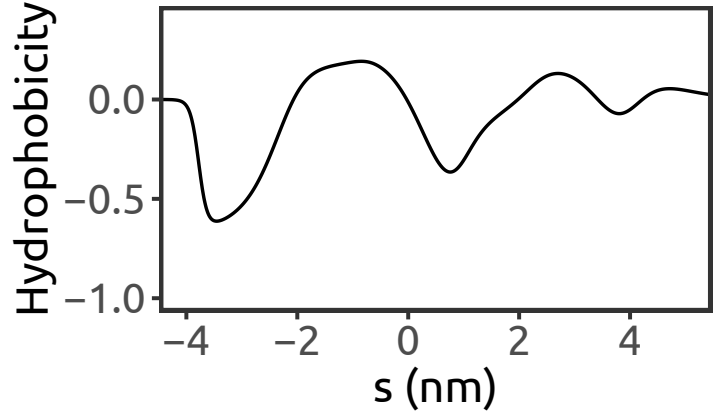
X-ray (2.2 Å)

Takeda et al., 2014

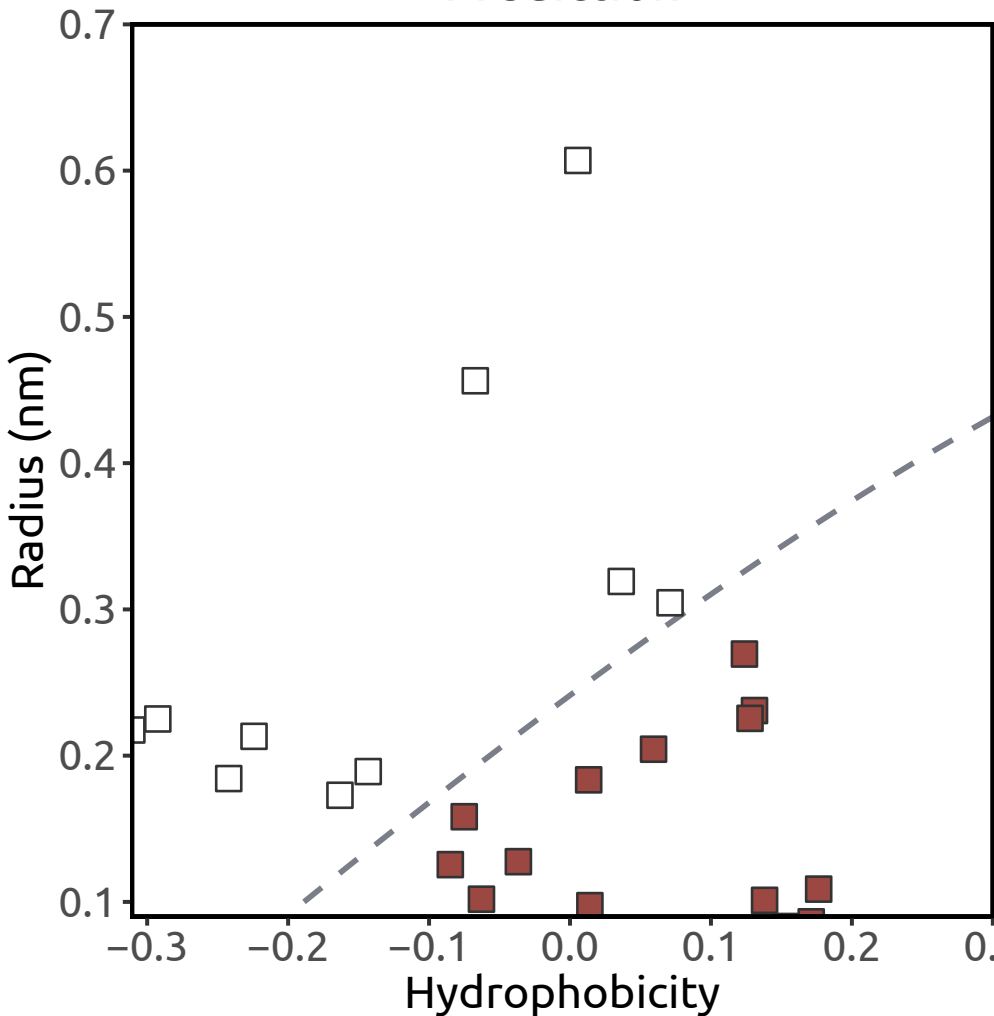
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.94 (n = 16)

Simulation result:

barrier to water

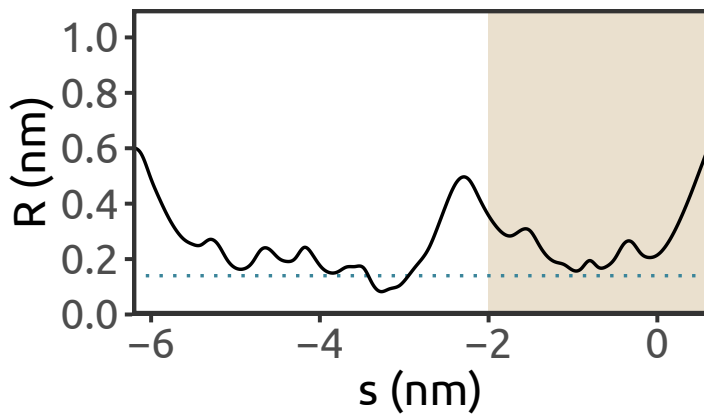
MscL (PDB ID: 2OAR)

Mycobacterium tuberculosis

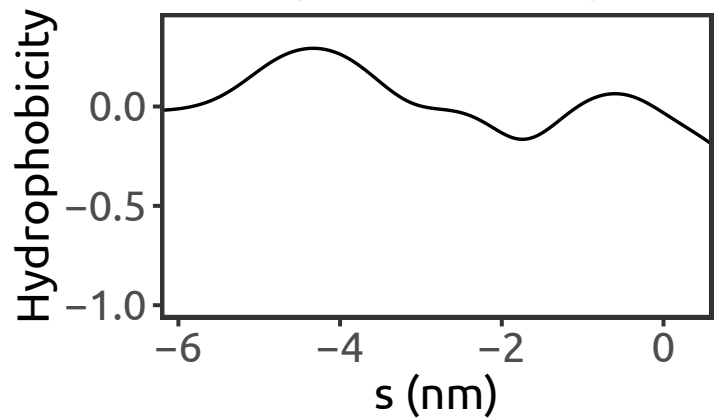
X-ray (3.5 Å)

Chang et al., 1998

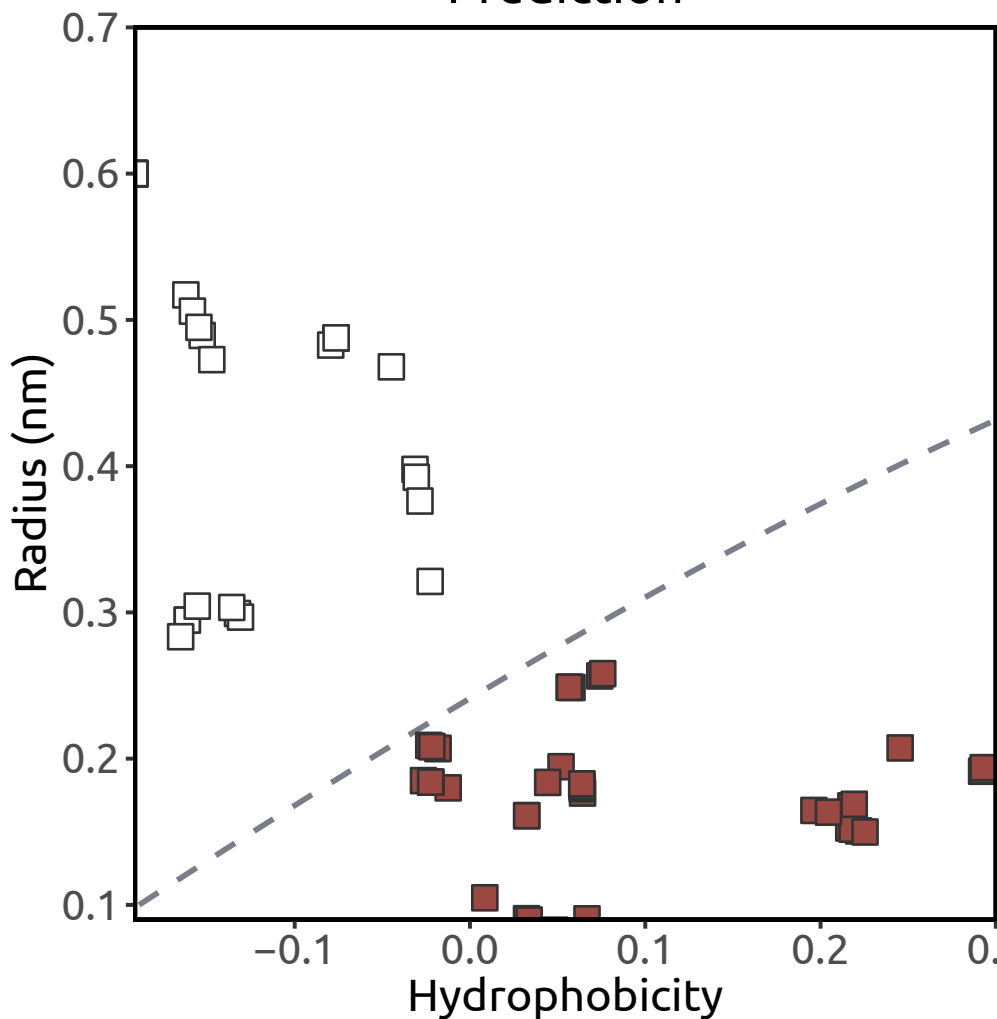
Pore radius



Hydrophobicity



Prediction



Heuristic score:
4.73 (n = 44)

Simulation result:
barrier to water

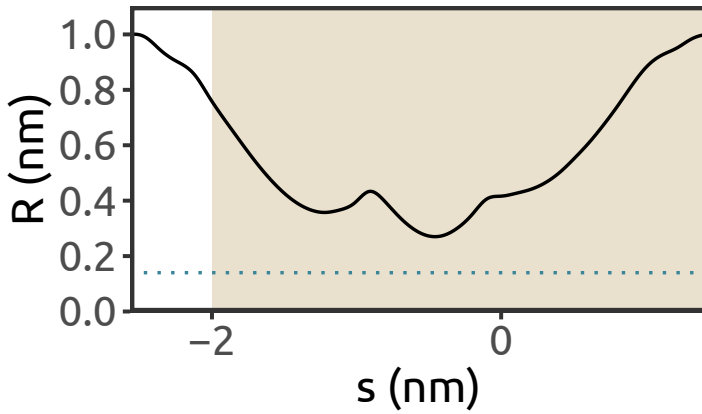
MscL (PDB ID: 3HZQ)

Staphylococcus aureus

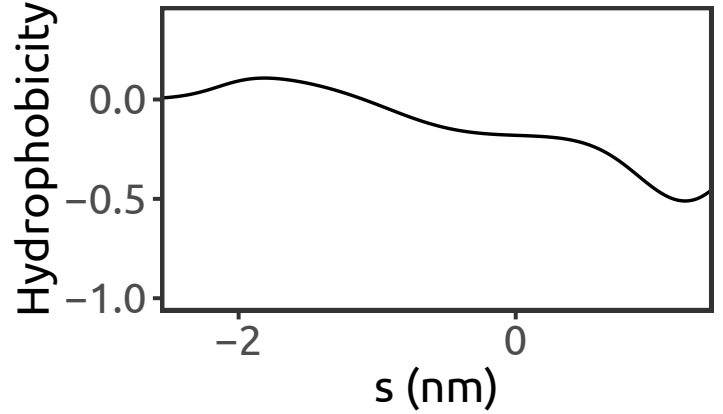
X-ray (3.82 Å)

Liu et al., 2009

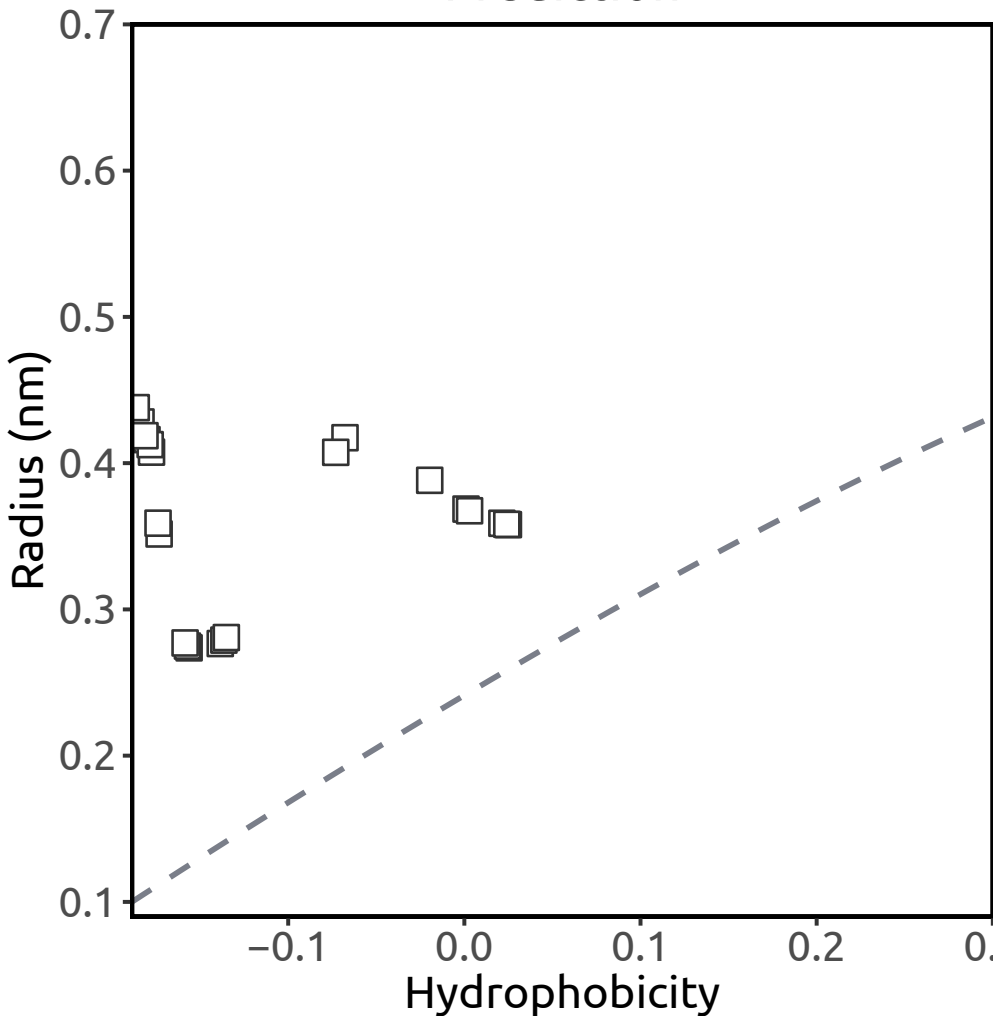
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 ($n = 0$)

Simulation result:
hydrated channel

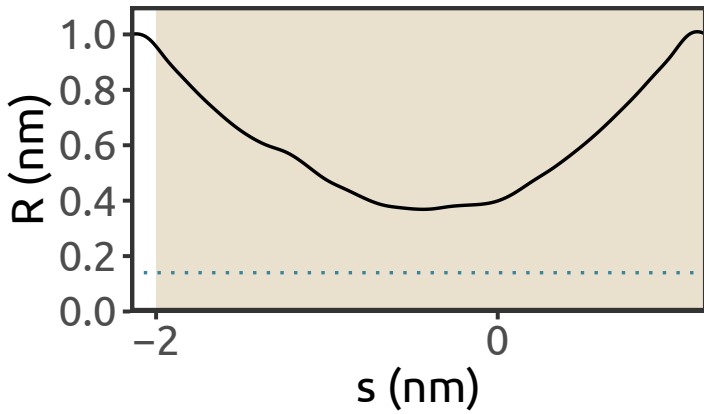
MscL (PDB ID: 4Y7J)

Methanosarcina acetivorans

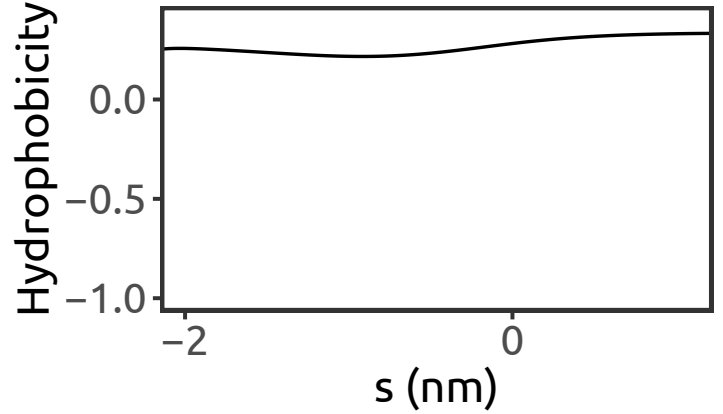
X-ray (4.1 Å)

Li et al., 2015

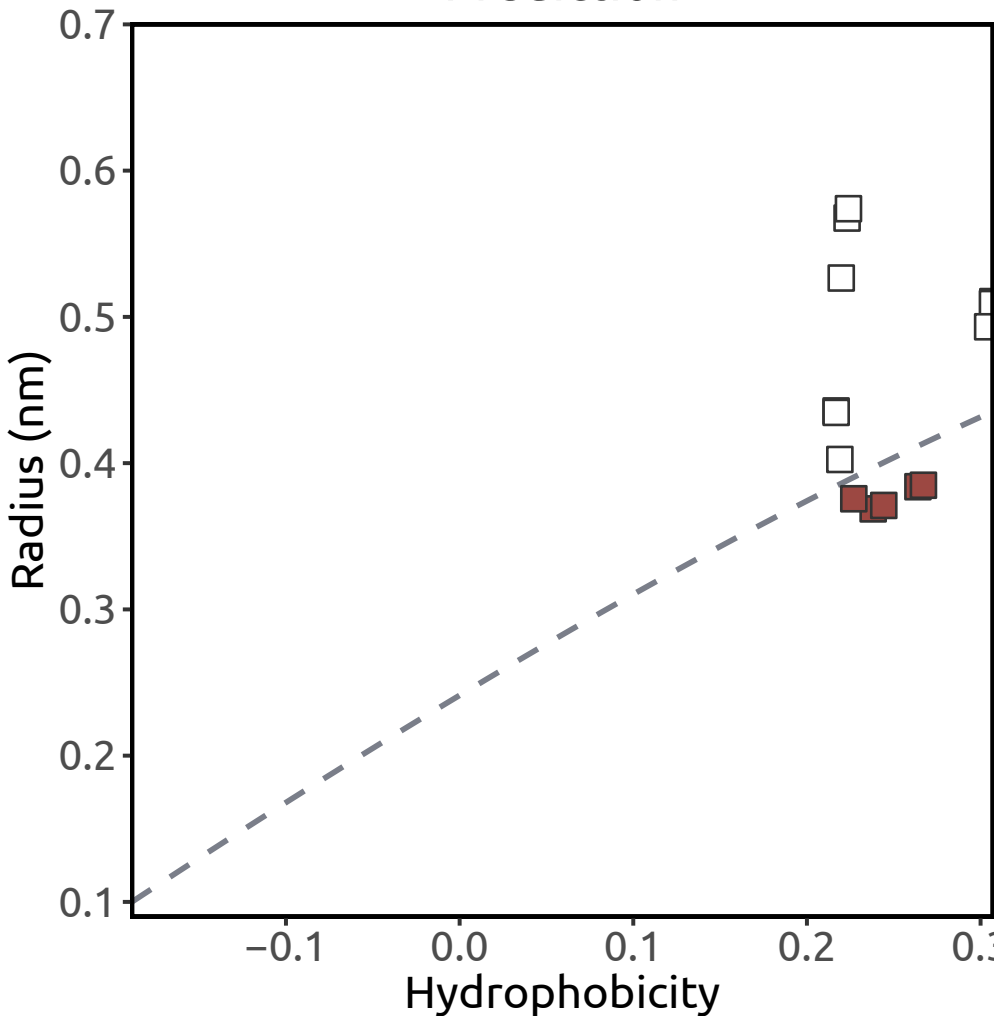
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.11 (n = 5)

Simulation result:
hydrated channel

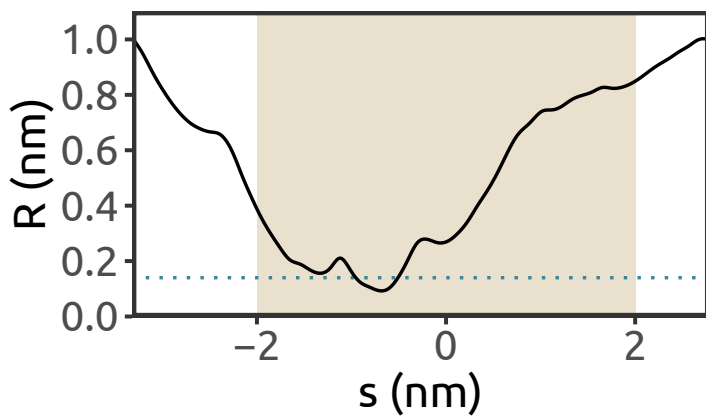
MscL (PDB ID: 4Y7K)

Methanosarcina acetivorans

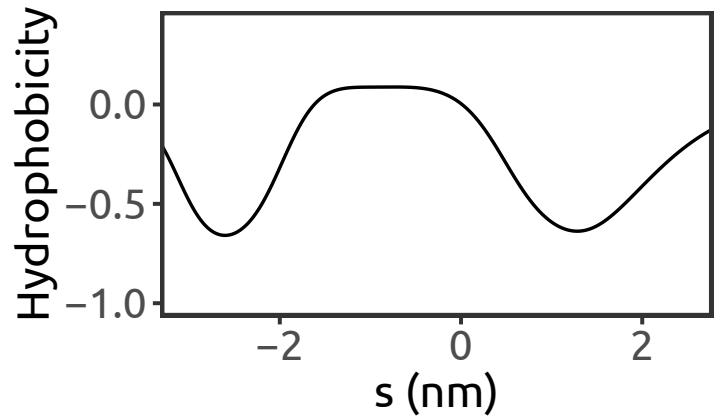
X-ray (3.5 Å)

Li et al., 2015

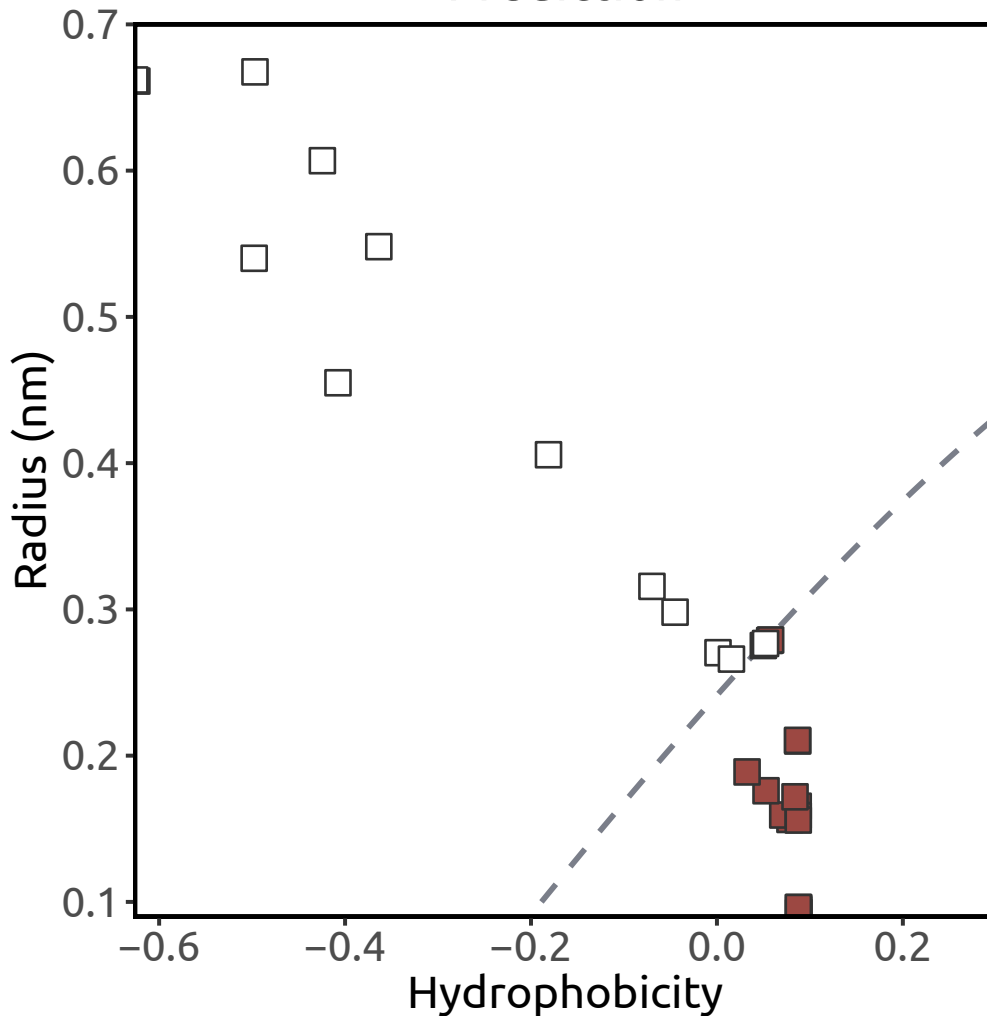
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.83 (n = 17)

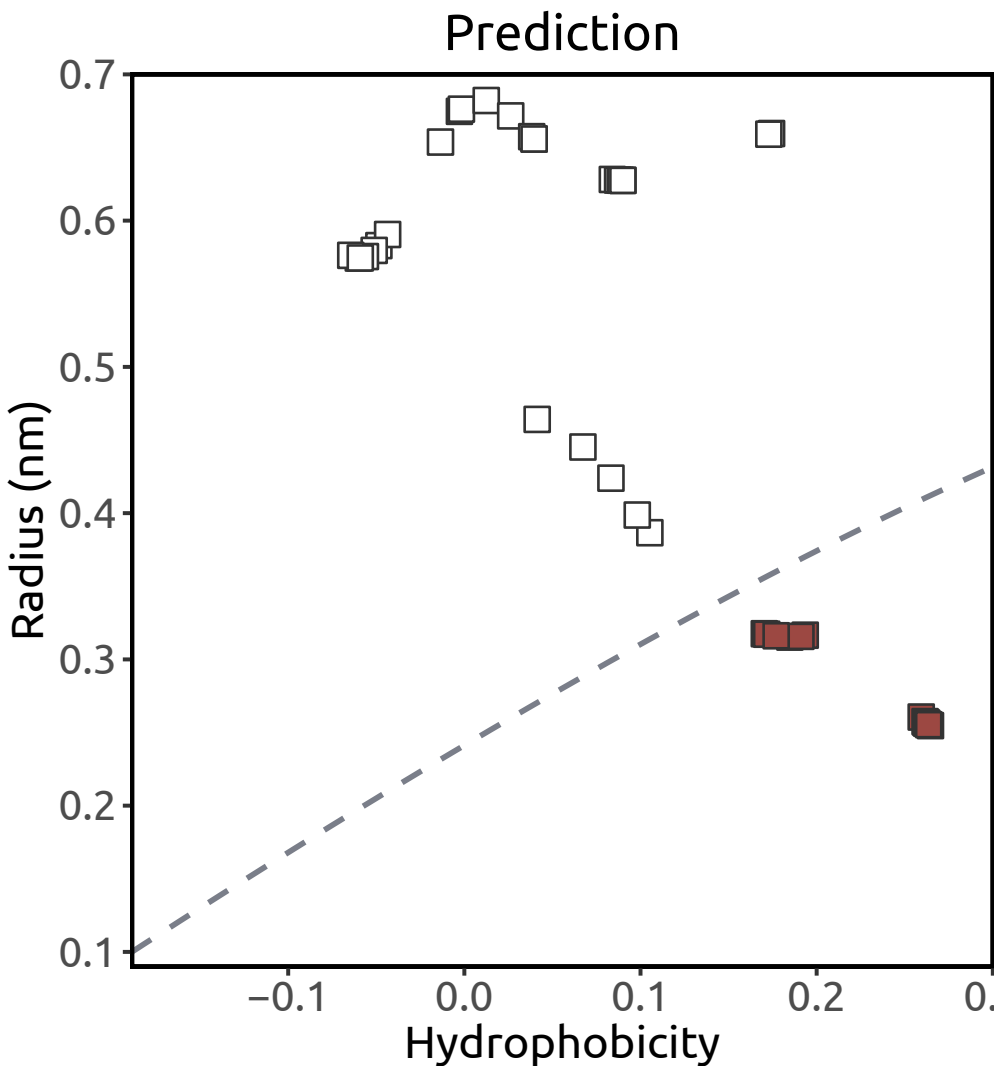
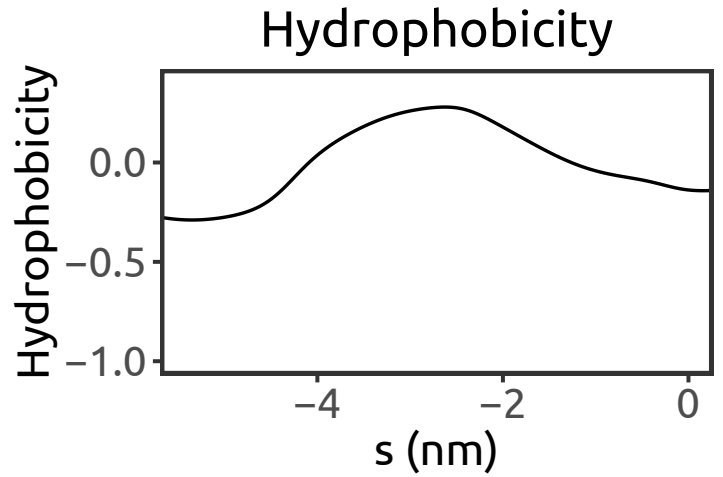
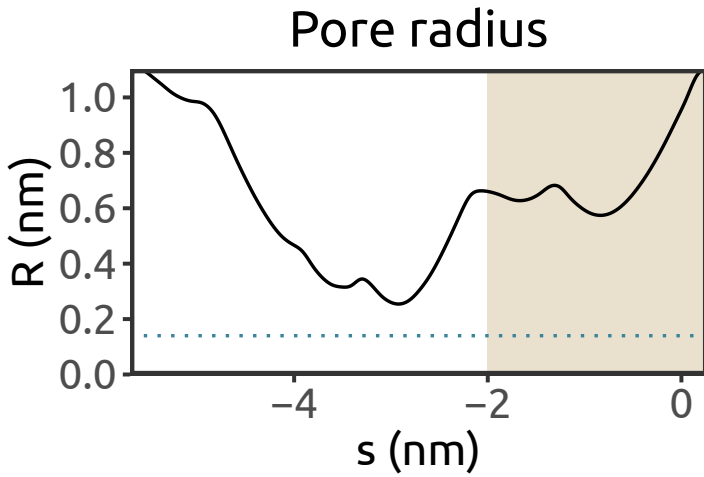
Simulation result:
barrier to water

MscS (PDB ID: 2OAU)

Escherichia coli

X-ray (3.7 Å)

Bass et al., 2002



Heuristic score:

1.03 (n = 12)

Simulation result:

barrier to water

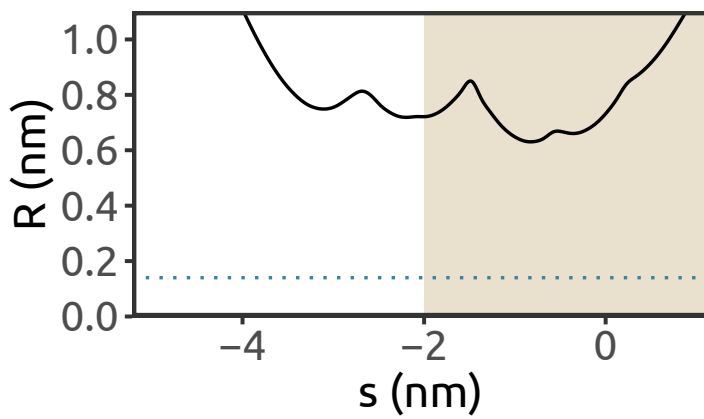
MscS (PDB ID: 2VV5)

Escherichia coli

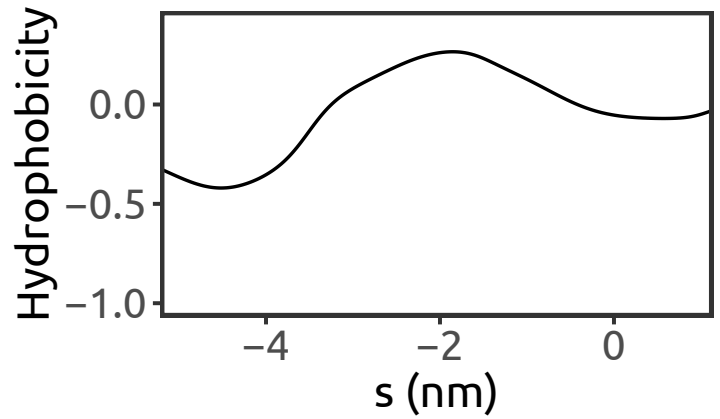
X-ray (3.45 Å)

Wang et al., 2008

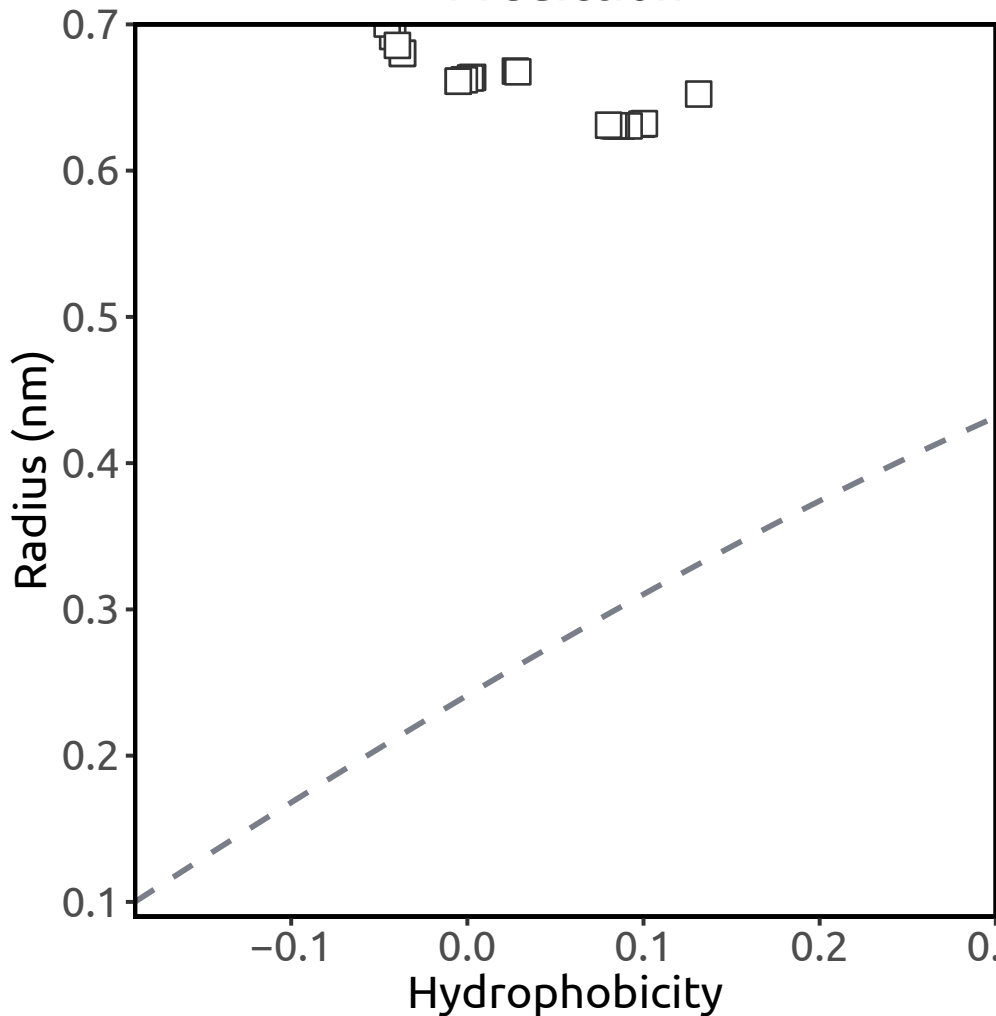
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

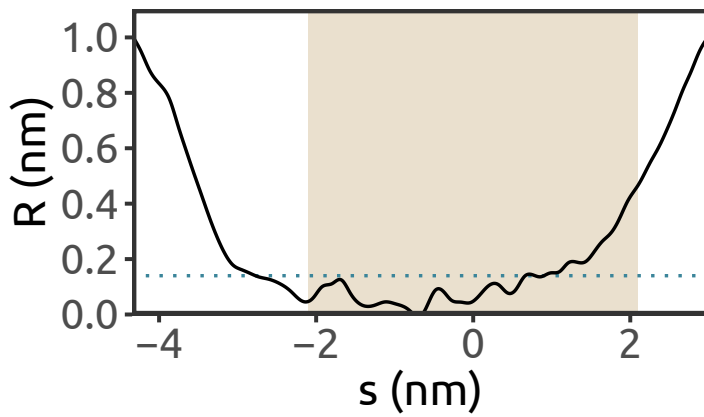
NNT (PDB ID: 5UNI)

Thermus thermophilus

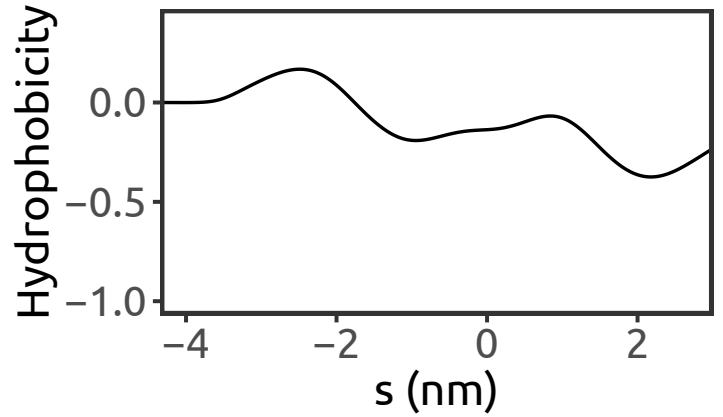
X-ray (2.2 Å)

Padayatti et al., 2017

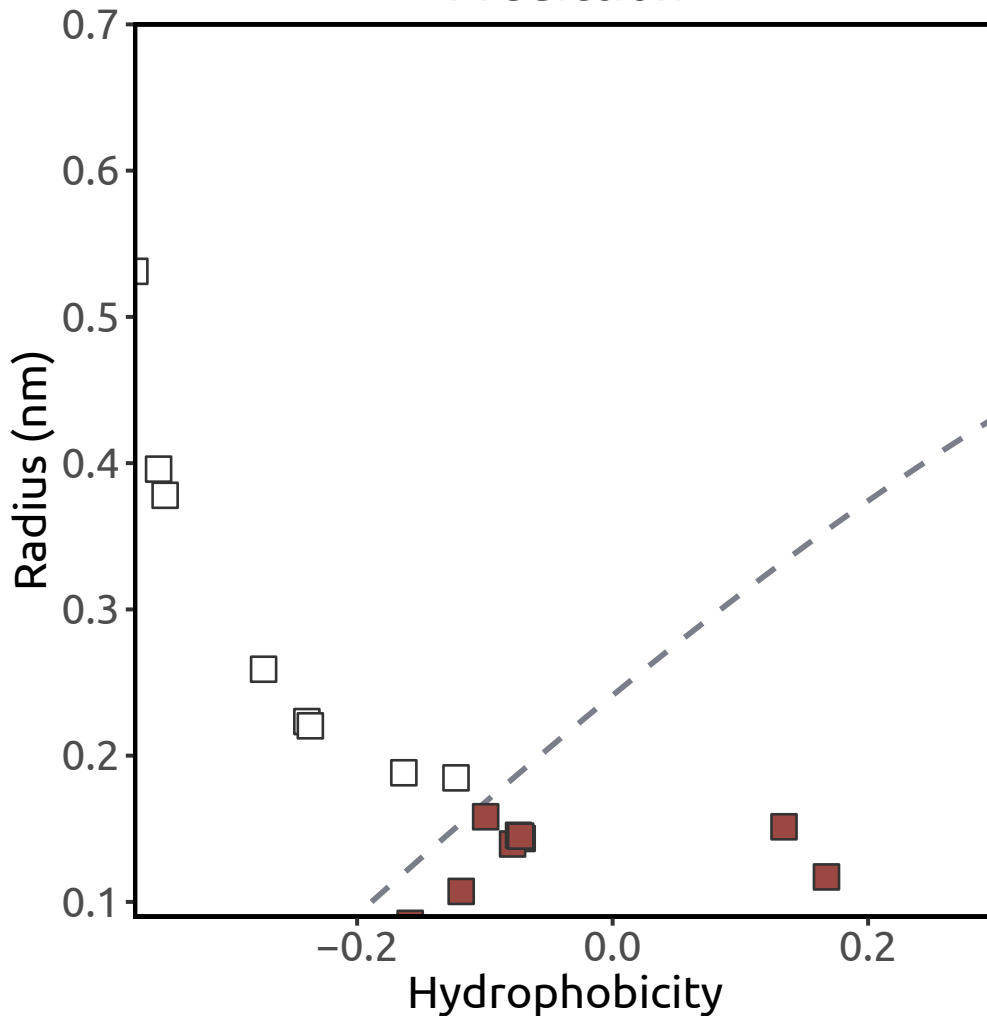
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.72 (n = 20)

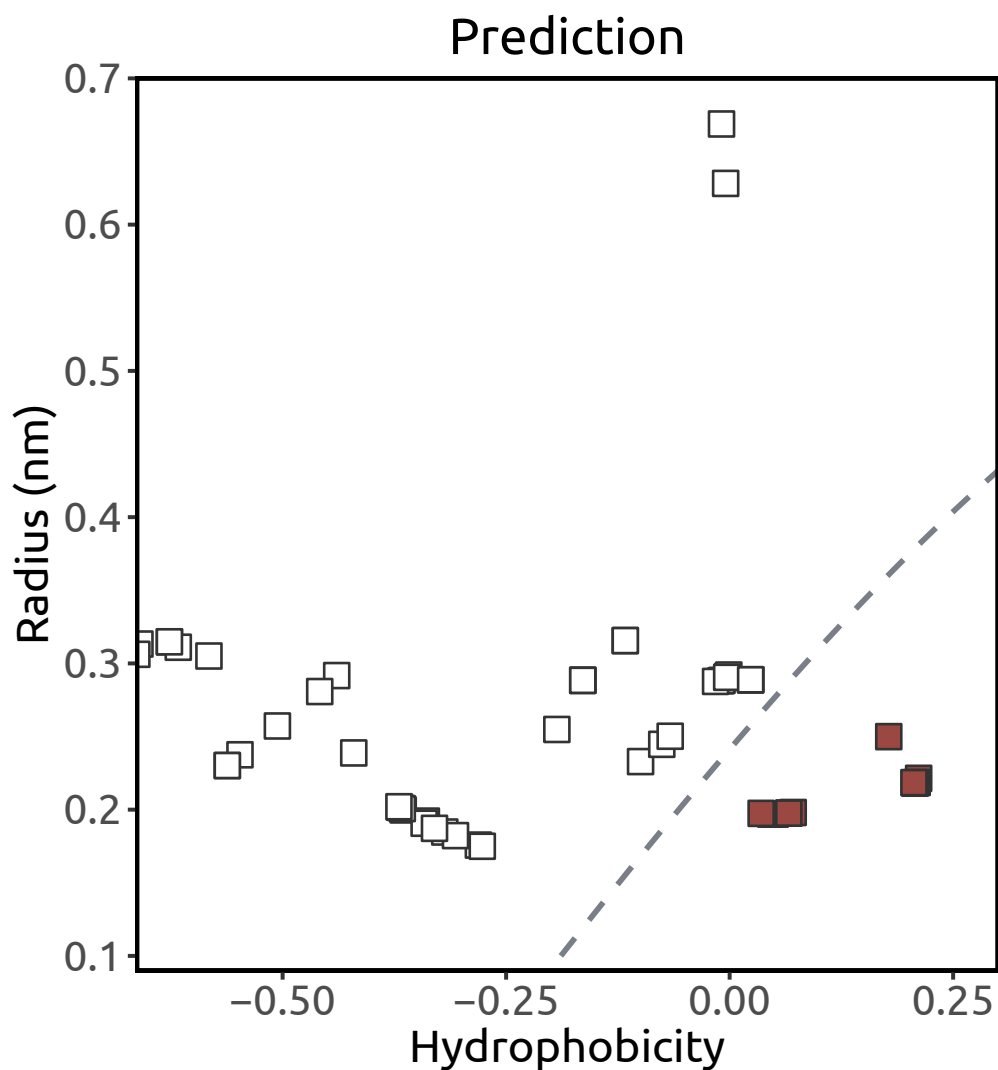
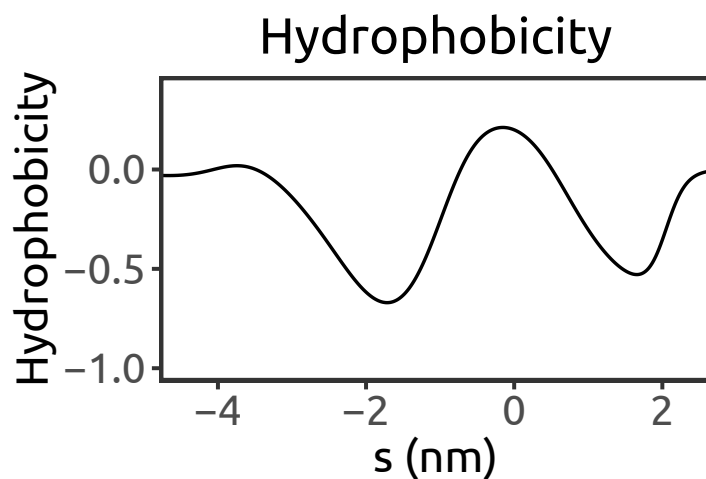
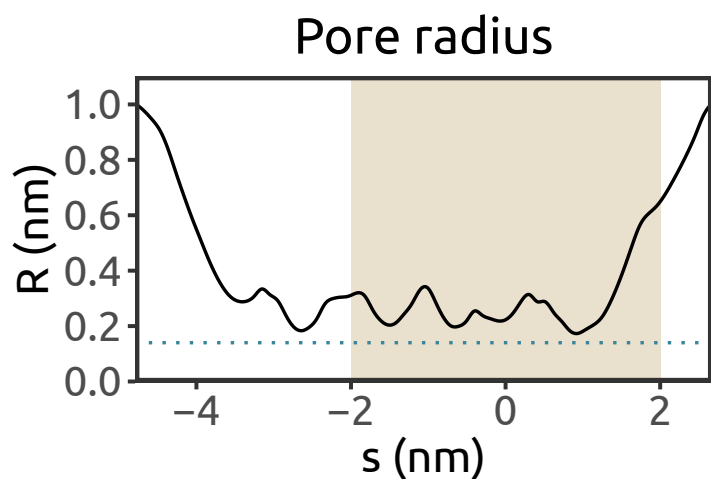
Simulation result:
barrier to water

Orai (PDB ID: 4HKR)

Drosophila melanogaster

X-ray (3.35 Å)

Hou et al., 2012



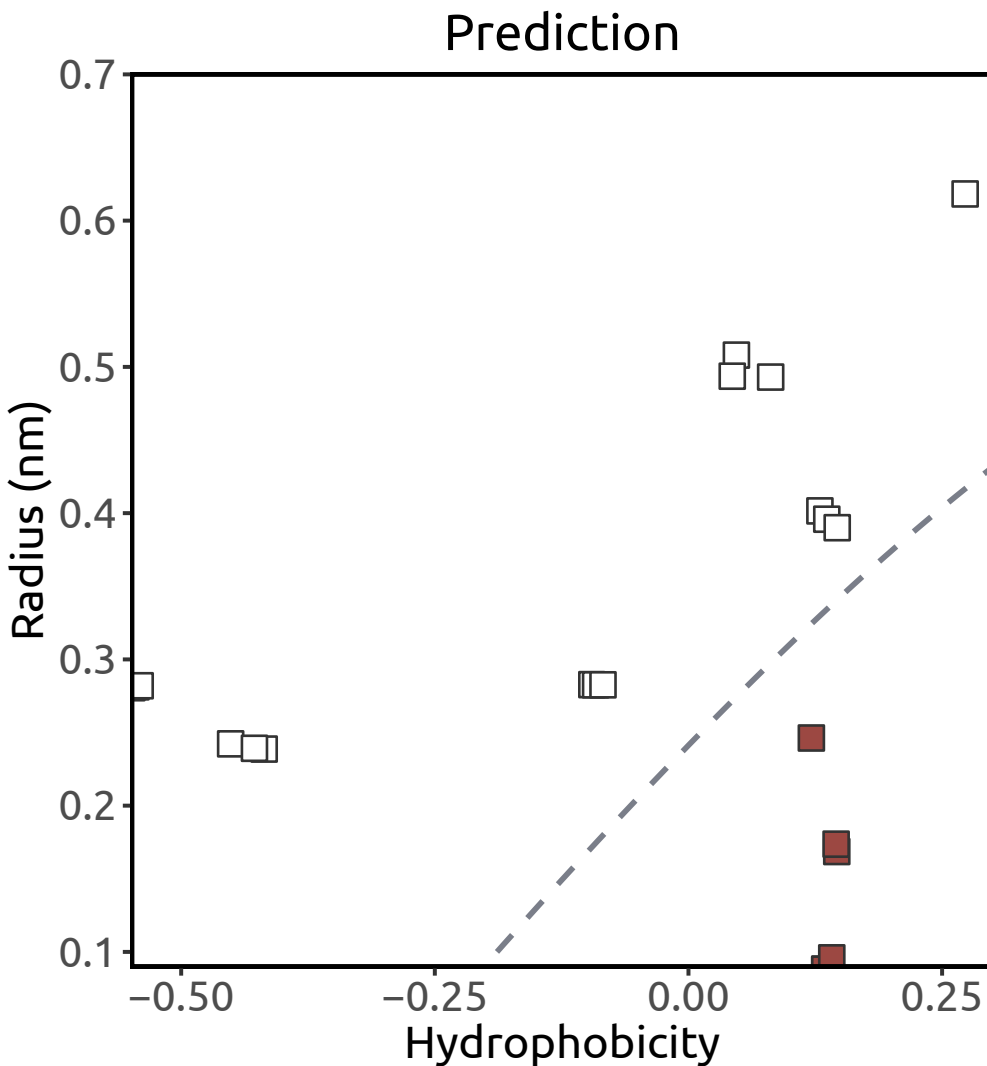
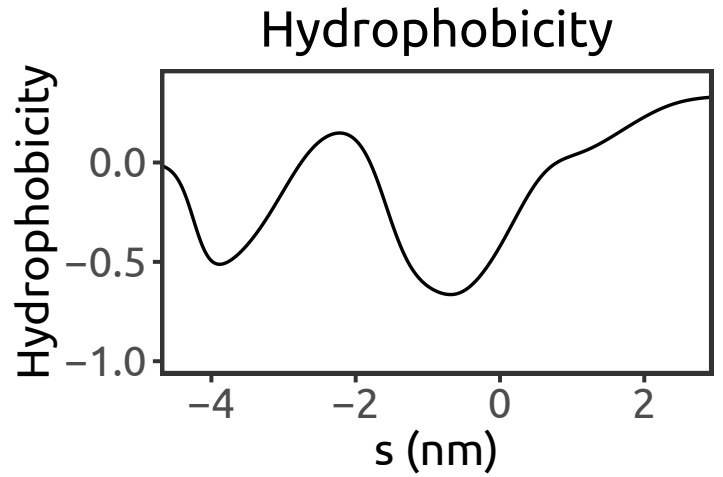
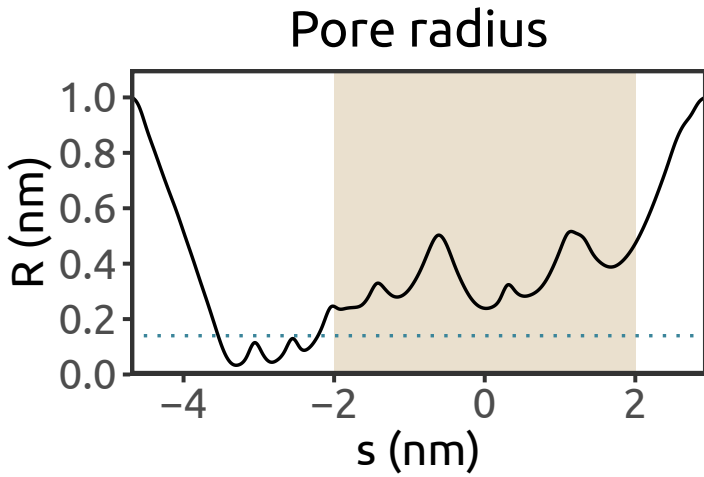
Heuristic score:
1.18 (n = 12)

Simulation result:
barrier to water

Piezo1 (PDB ID: 6BPZ)

Mus musculus
cryo-EM (3.8 Å)

Saotome et al., 2017



Heuristic score:
1.33 (n = 9)

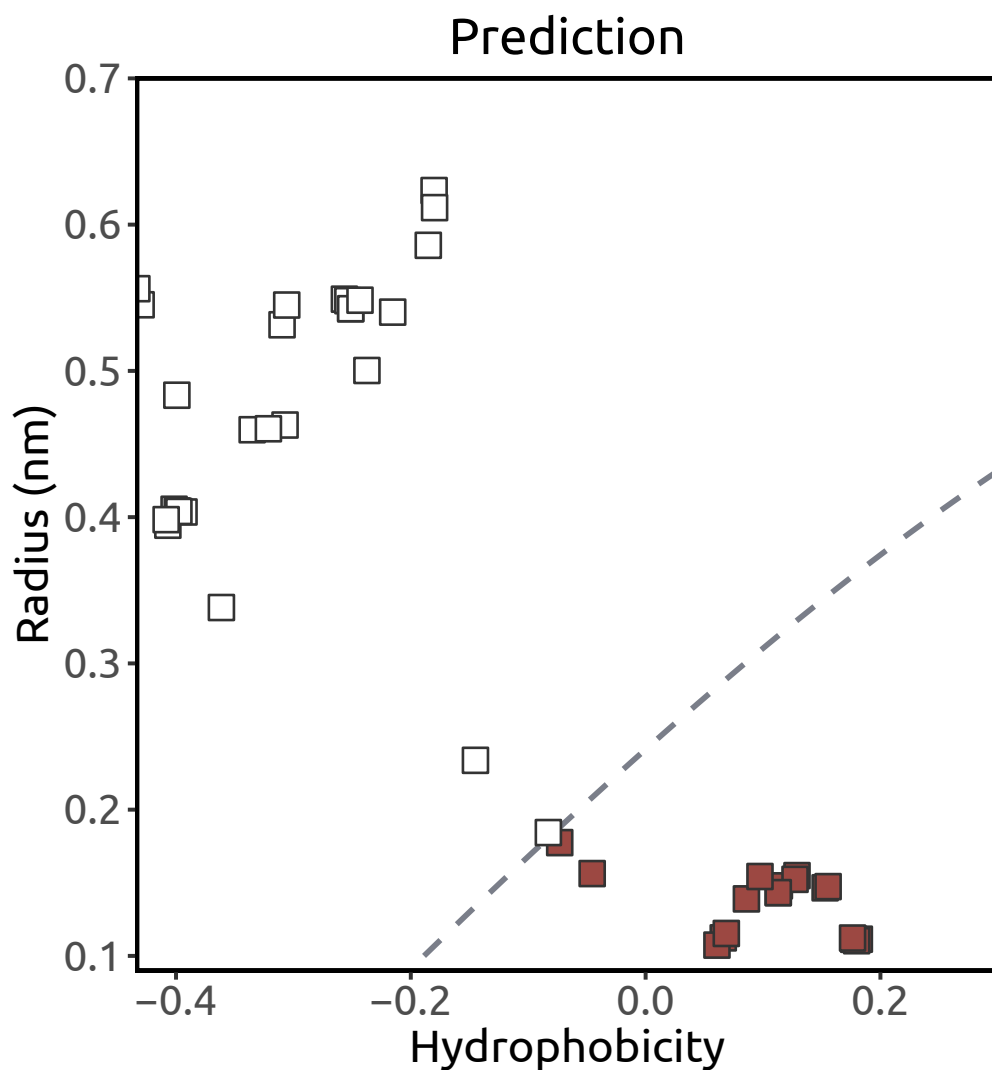
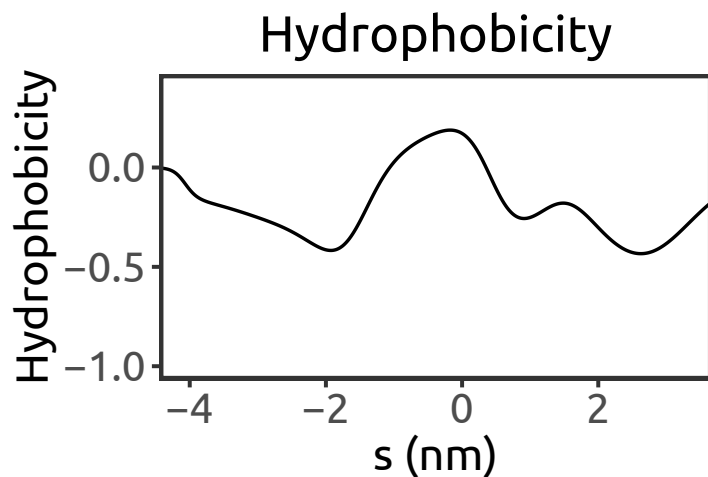
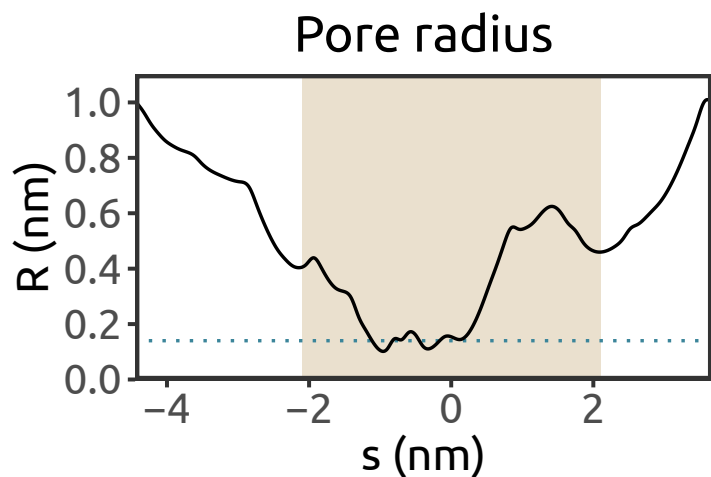
Simulation result:
barrier to water

TMEM175 (PDB ID: 5VRE)

Chamaesiphon minutus

X-ray (3.30 Å)

Lee et al., 2017



Heuristic score:
2.63 (n = 18)

Simulation result:
barrier to water

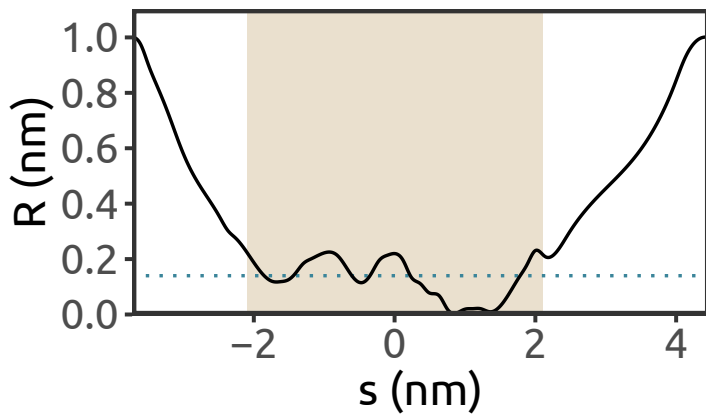
TRIC (PDB ID: 5EGI)

Caenorhabditis elegans

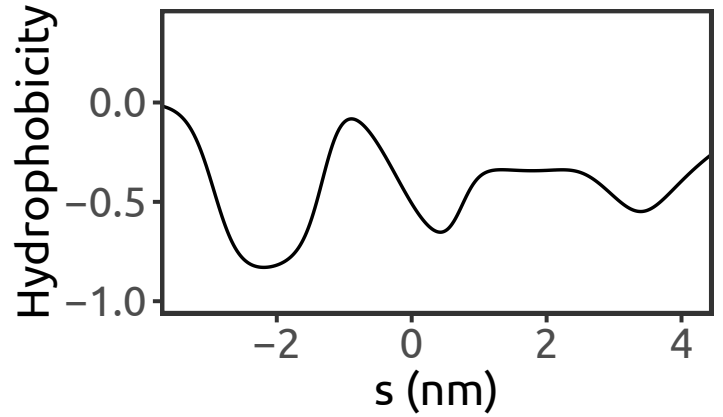
X-ray (3.3 Å)

Yang et al., 2016

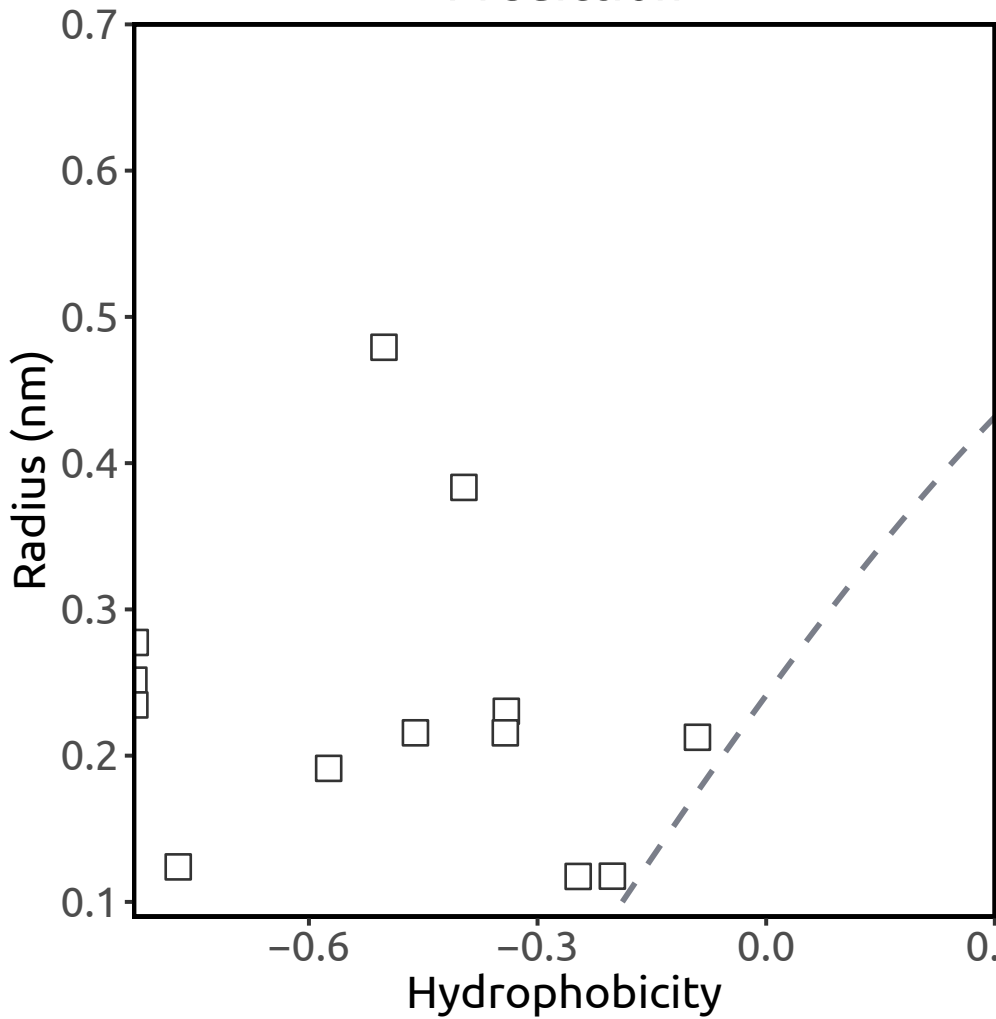
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
barrier to water

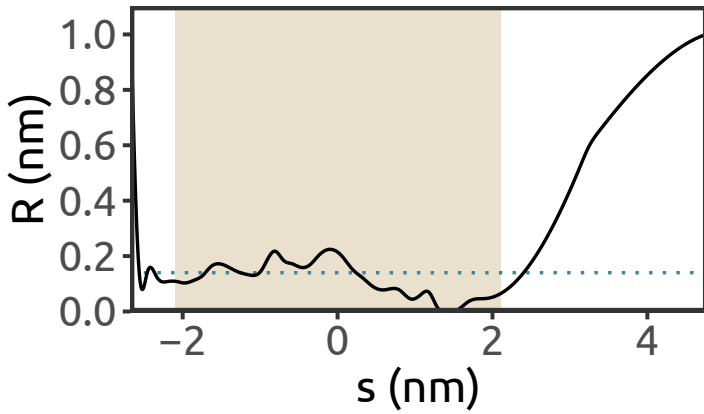
TRIC (PDB ID: 5EIK)

Caenorhabditis elegans

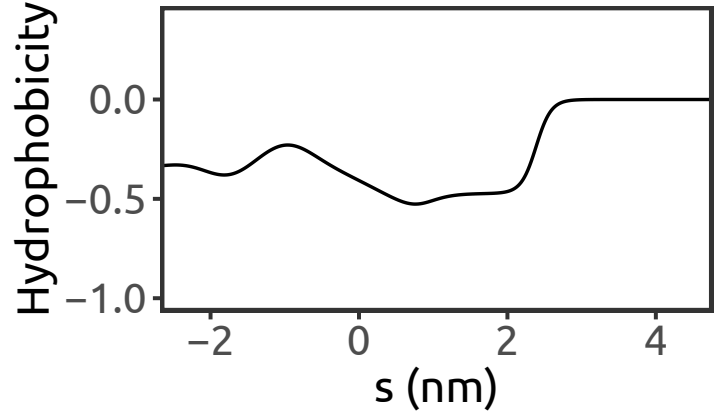
X-ray (2.3 Å)

Yang et al., 2016

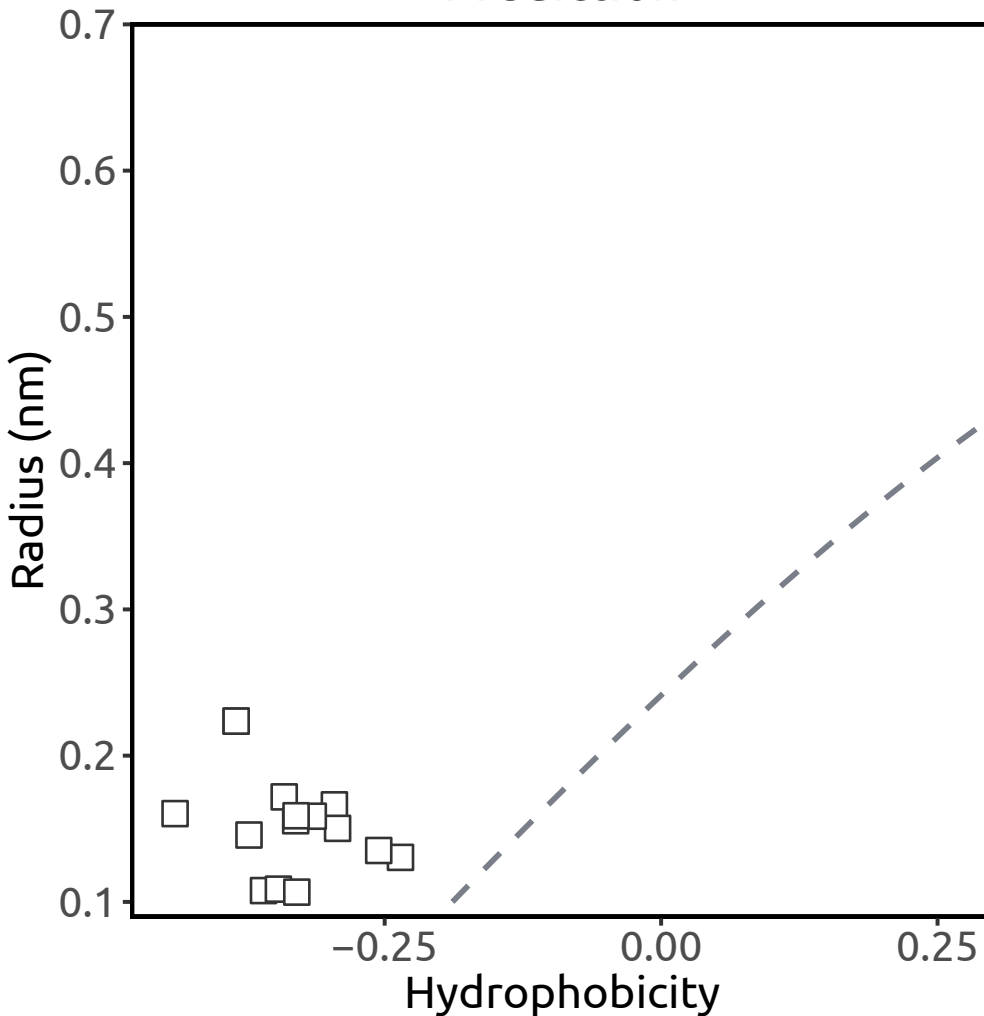
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
barrier to water

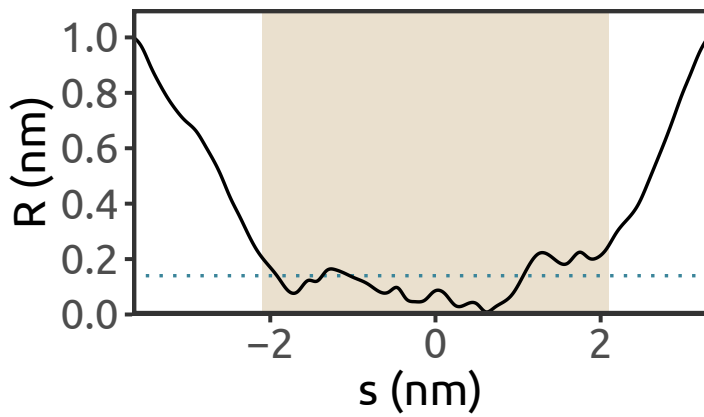
TRIC (PDB ID: 5WUC)

Sulfolobus acidocaldarius

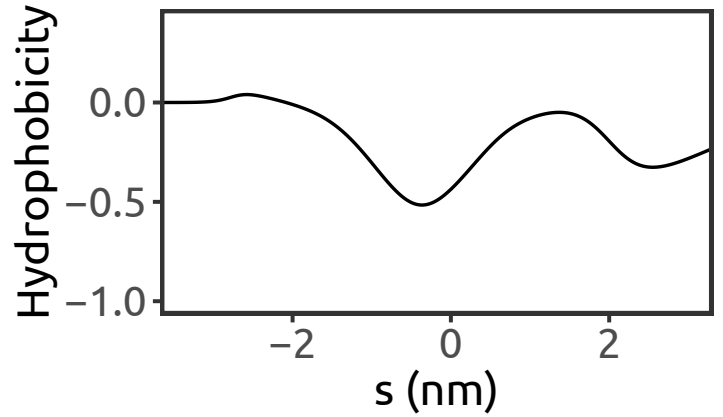
X-ray (1.6 Å)

Su et al., 2017

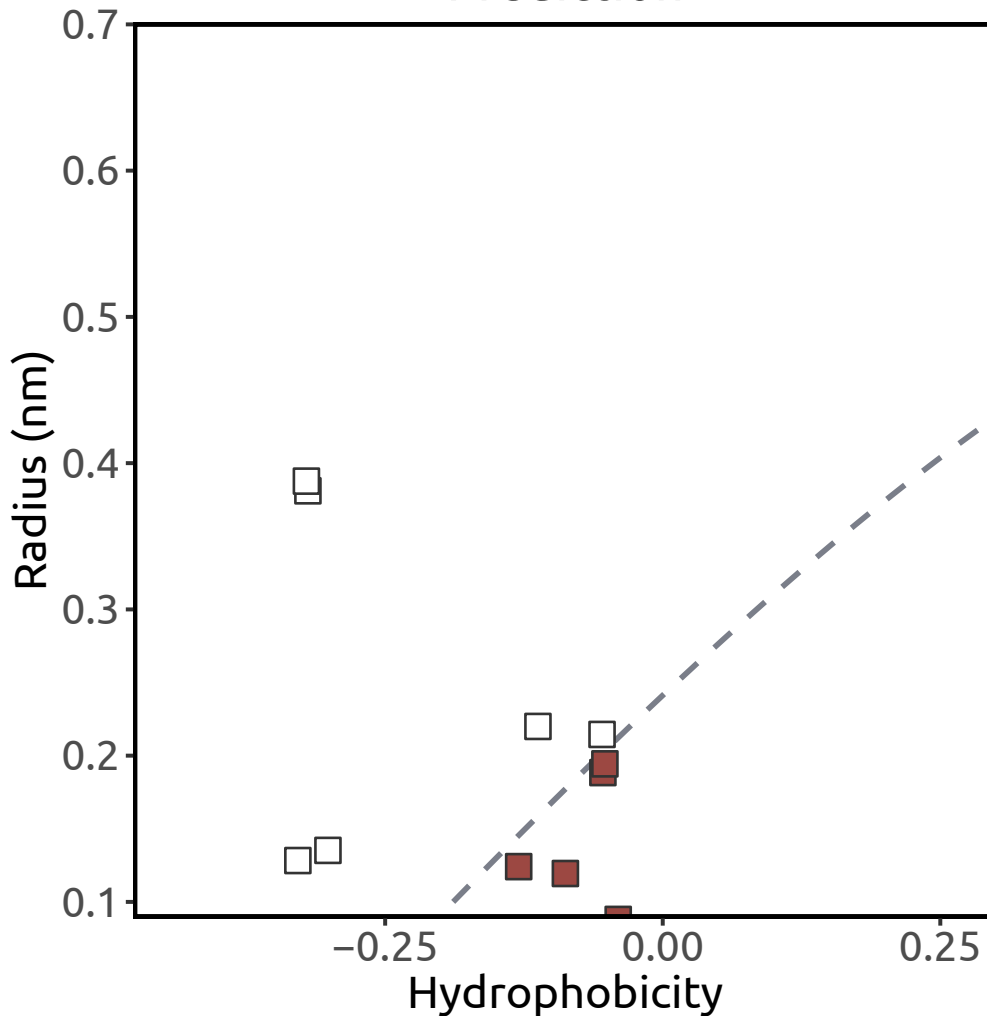
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.27 (n = 6)

Simulation result:
barrier to water

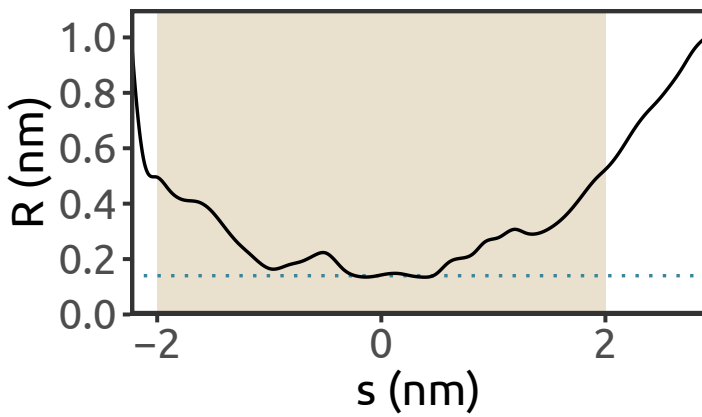
TRIC (PDB ID: 5WUE)

Sulfolobus acidocaldarius

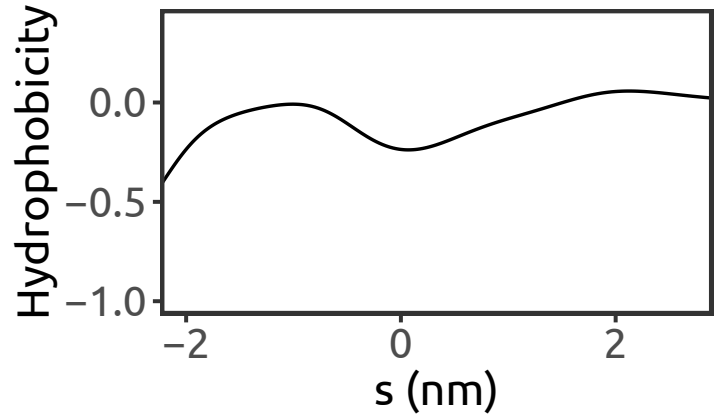
X-ray (2.4 Å)

Su et al., 2017

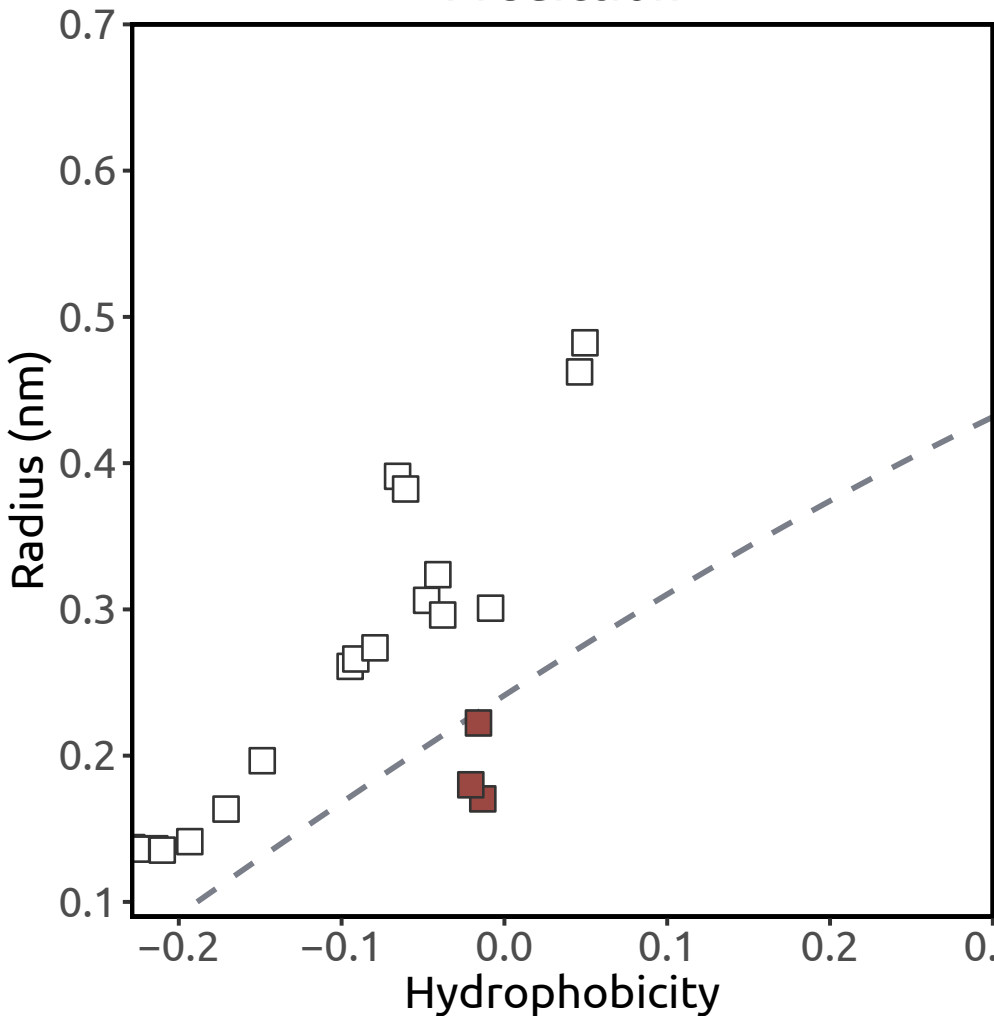
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.09 (n = 3)

Simulation result:
hydrated channel

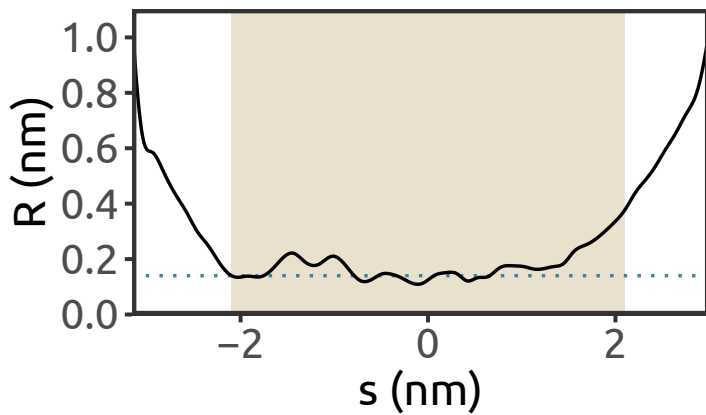
TRIC (PDB ID: 5WUF)

Colwellia psychrerythraea

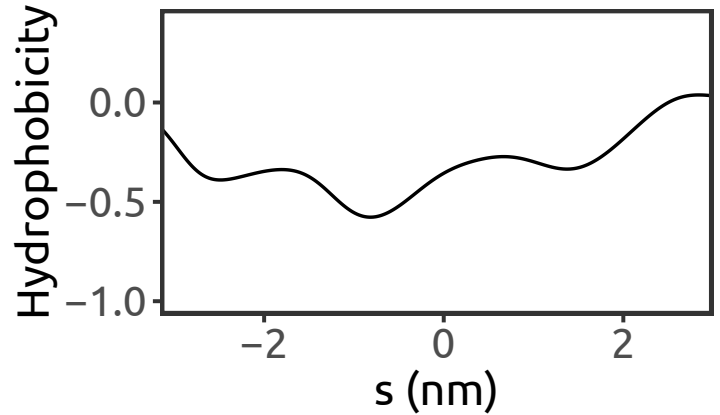
X-ray (2.40 Å)

Su et al., 2017

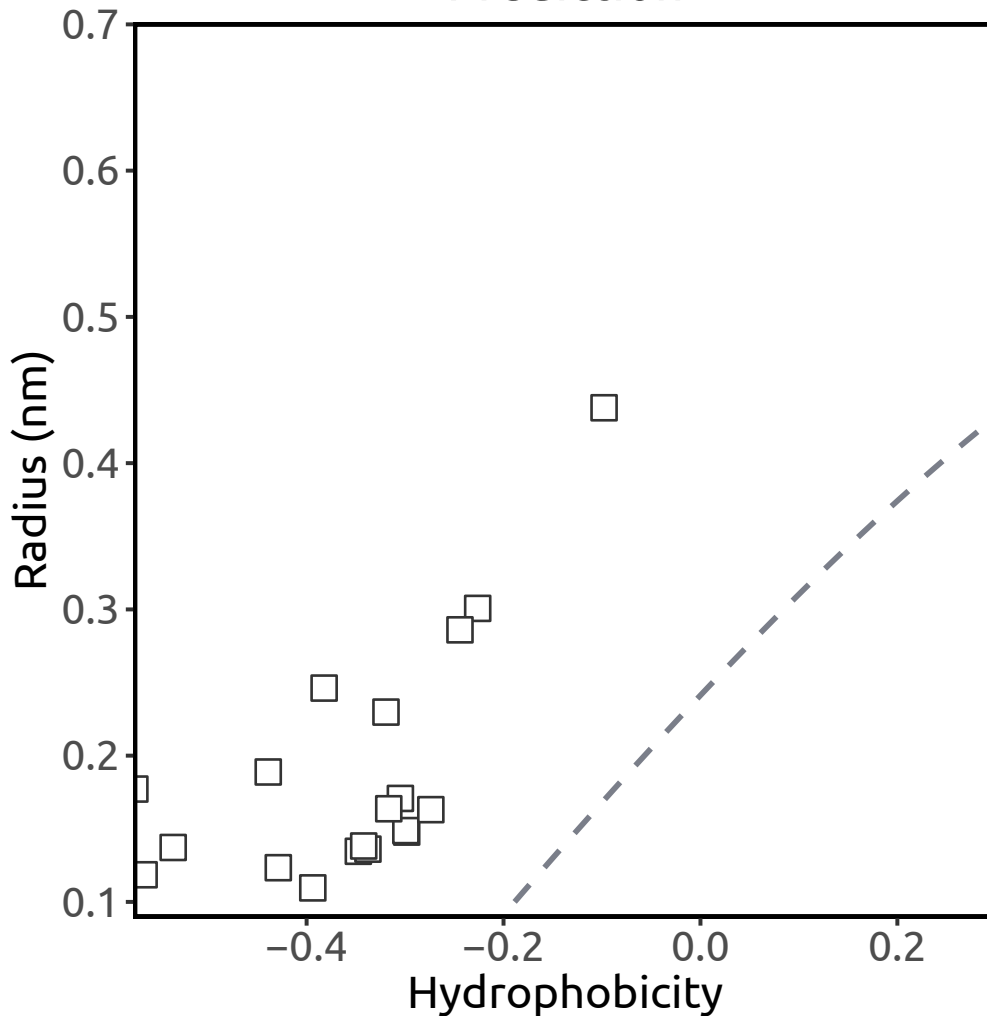
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0 (n = 0)

Simulation result:
hydrated channel

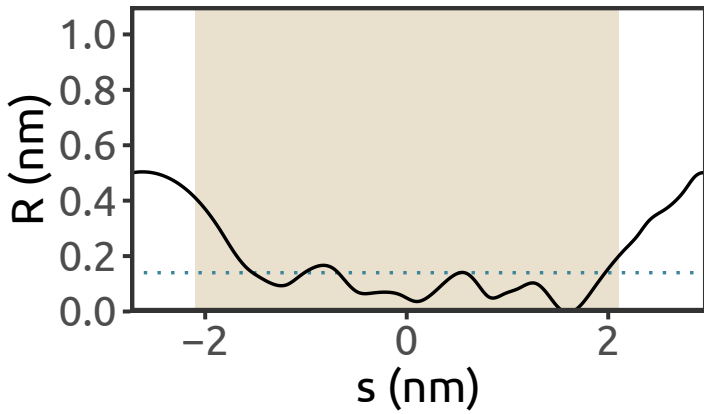
YetJ (PDB ID: 4PGR)

Bacillus subtilis

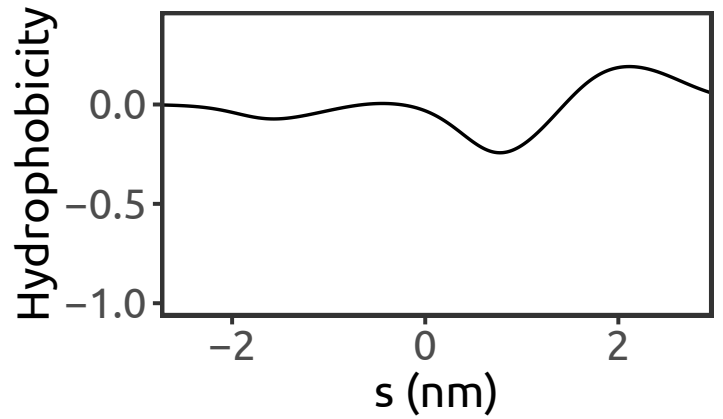
X-ray (1.95 Å)

Chang et al., 2014

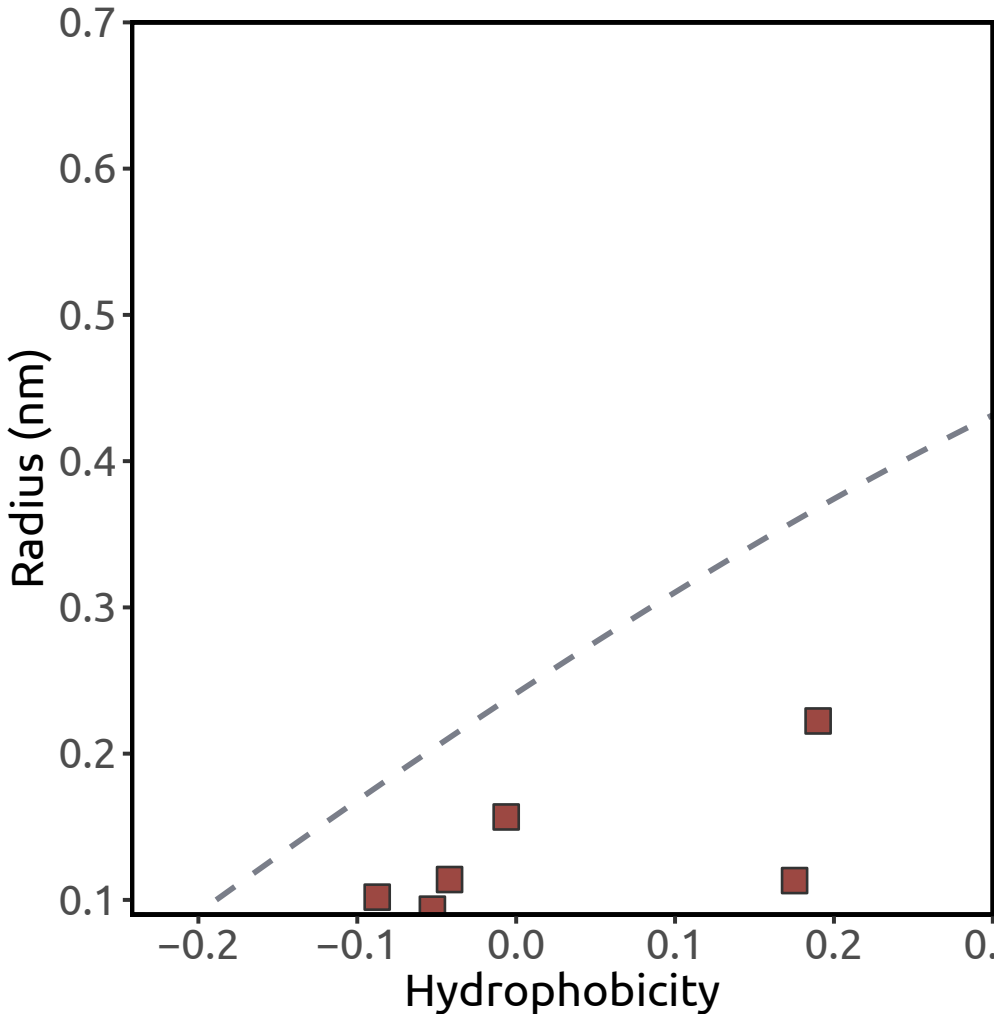
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.18 (n = 11)

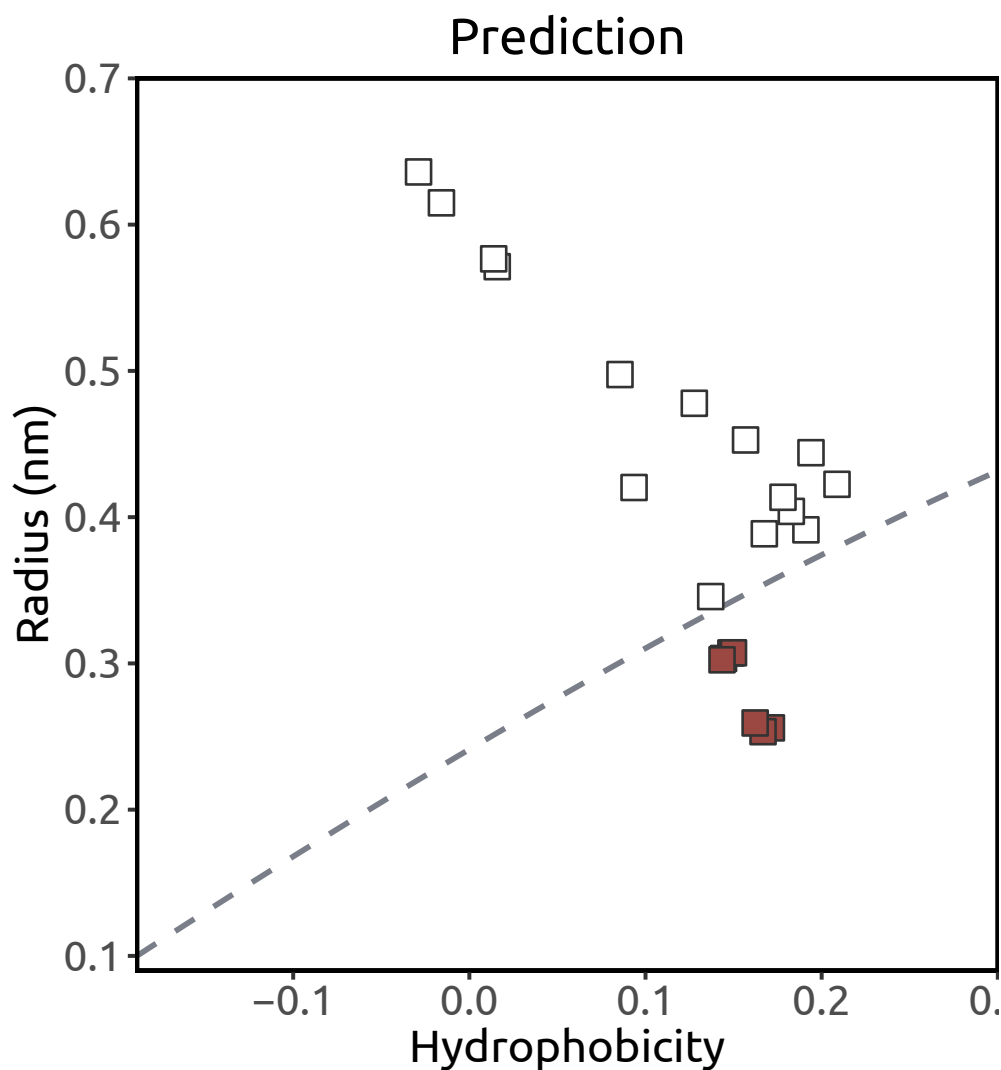
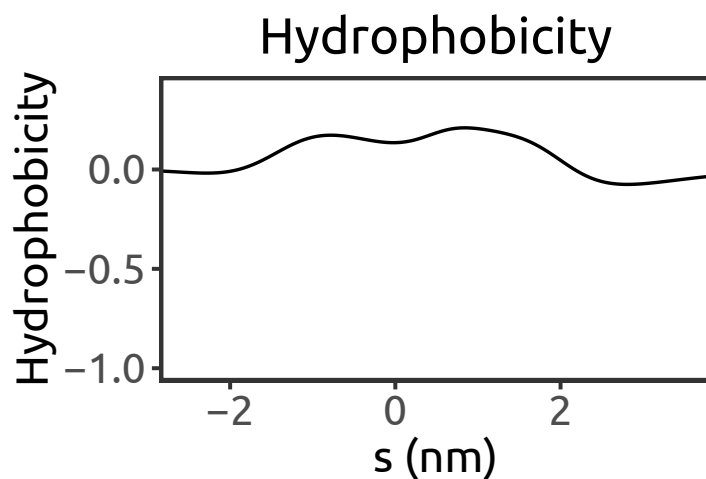
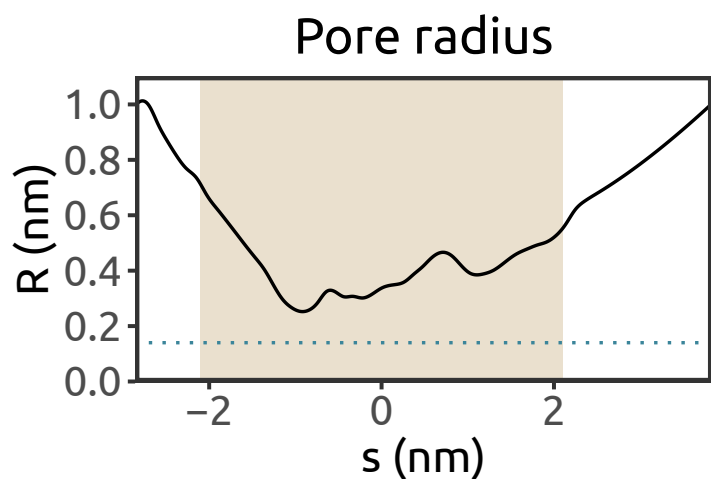
Simulation result:
barrier to water

YetJ (PDB ID: 4PGS)

Bacillus subtilis

X-ray (2.5 Å)

Chang et al., 2014



Heuristic score:
0.37 (n = 7)

Simulation result:
barrier to water

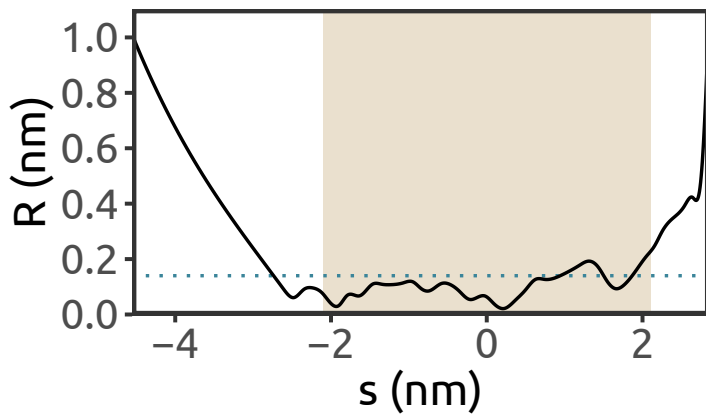
YetJ (PDB ID: 4PGU)

Bacillus subtilis

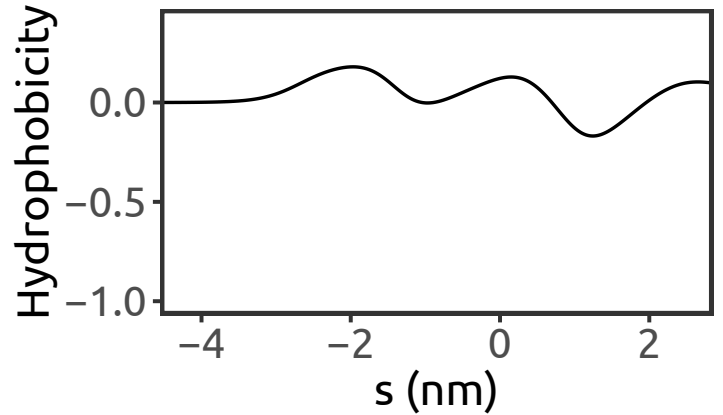
X-ray (3.40 Å)

Chang et al., 2014

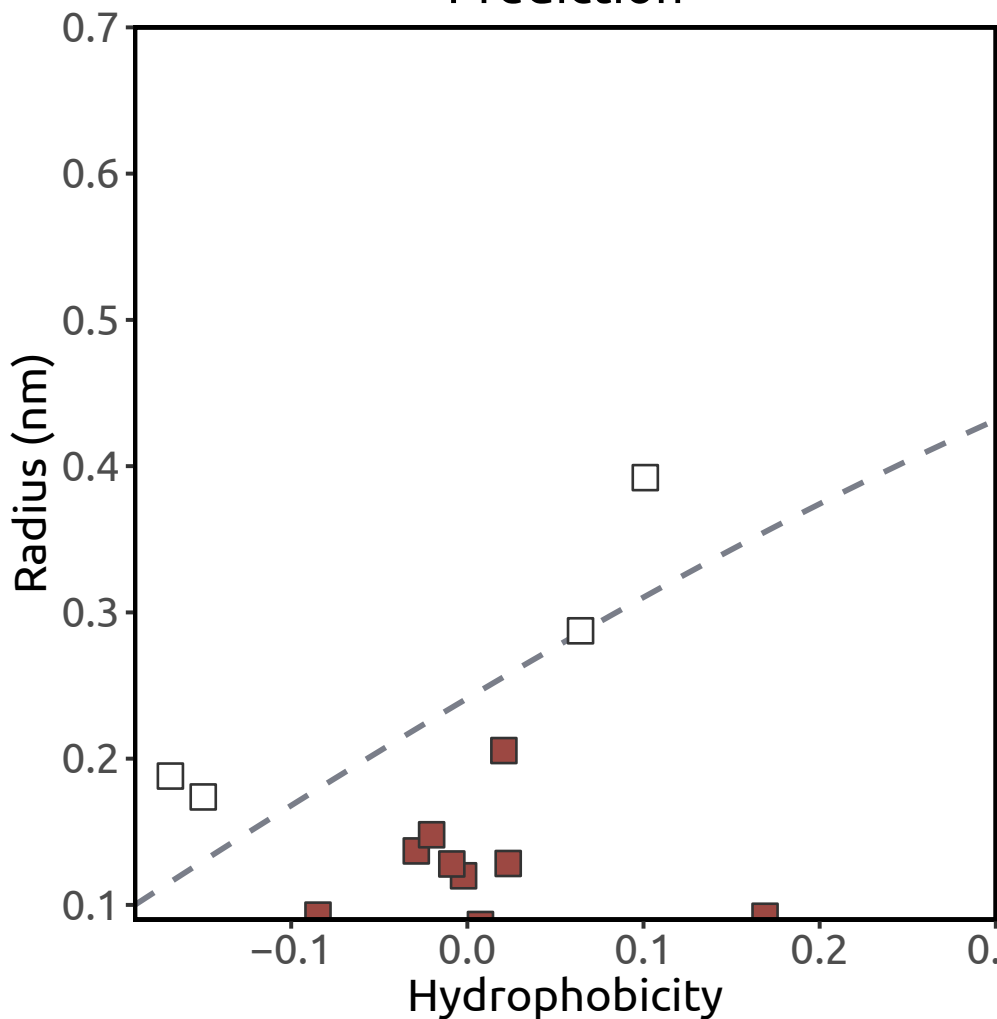
Pore radius



Hydrophobicity



Prediction



Heuristic score:

2 (n = 14)

Simulation result:

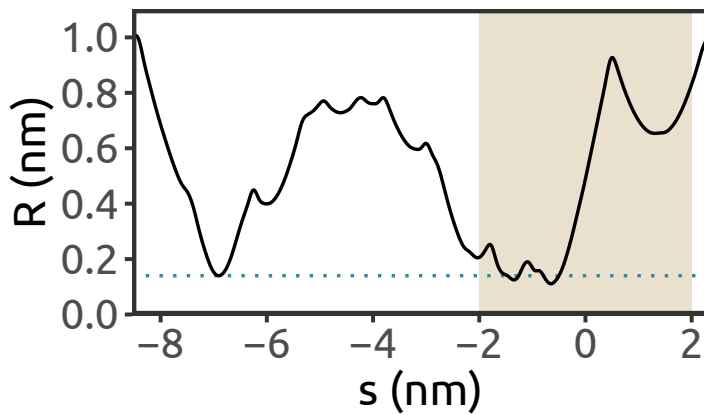
barrier to water

BEST1 (PDB ID: 4RDQ)

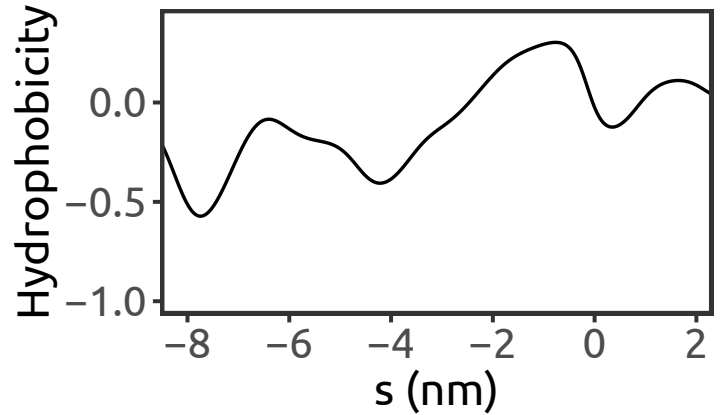
Gallus gallus
X-ray (2.85 Å)

Kane et al., 2014

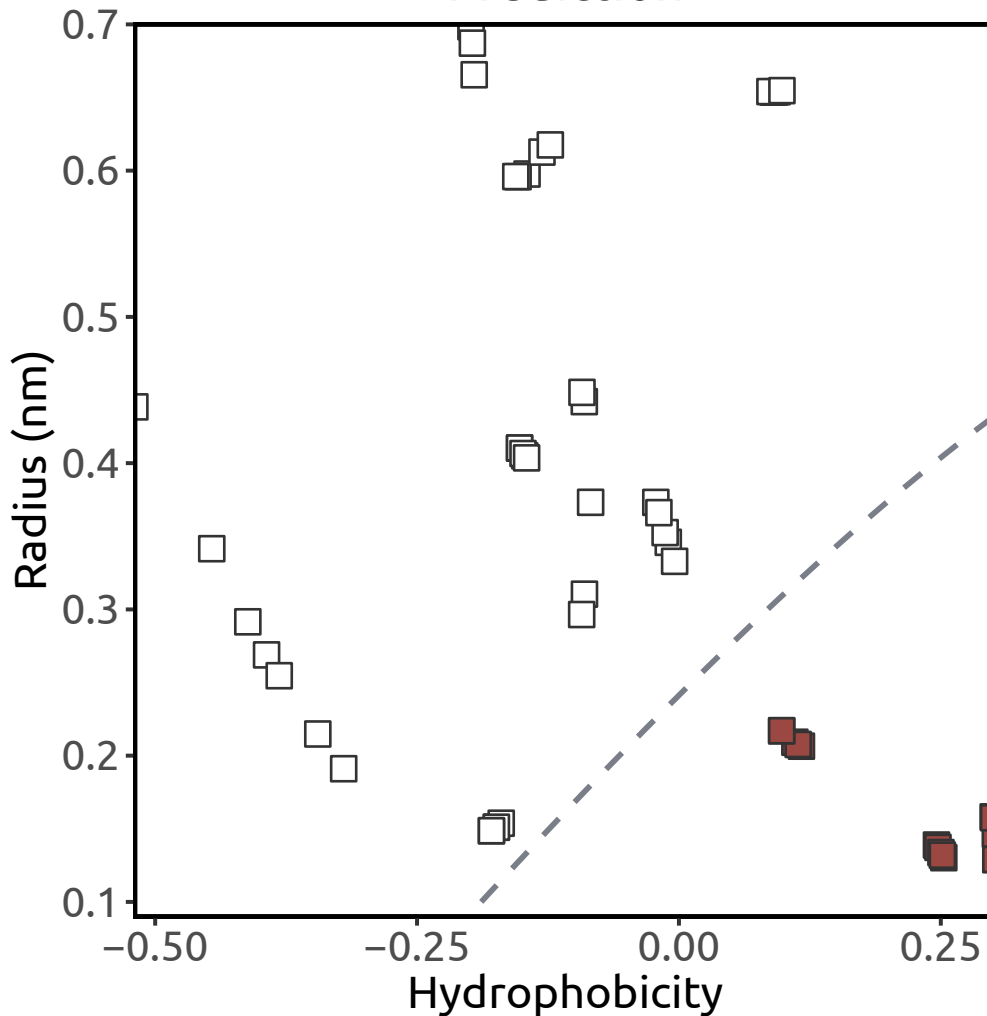
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.82 (n = 15)

Simulation result:
barrier to water

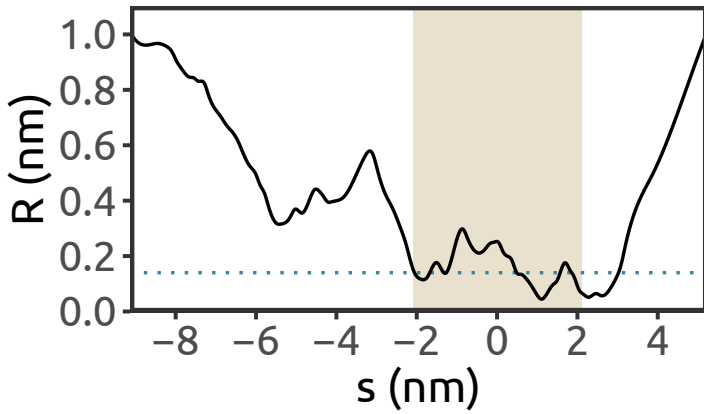
CFTR (PDB ID: 5UAK)

Homo sapiens

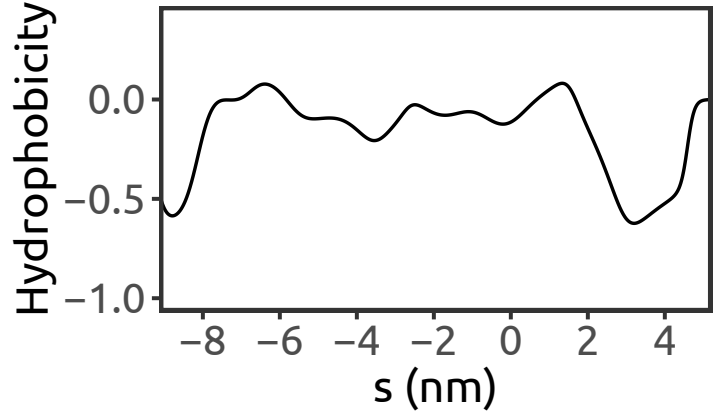
cryo-EM (3.87 Å)

Liu et al., 2017

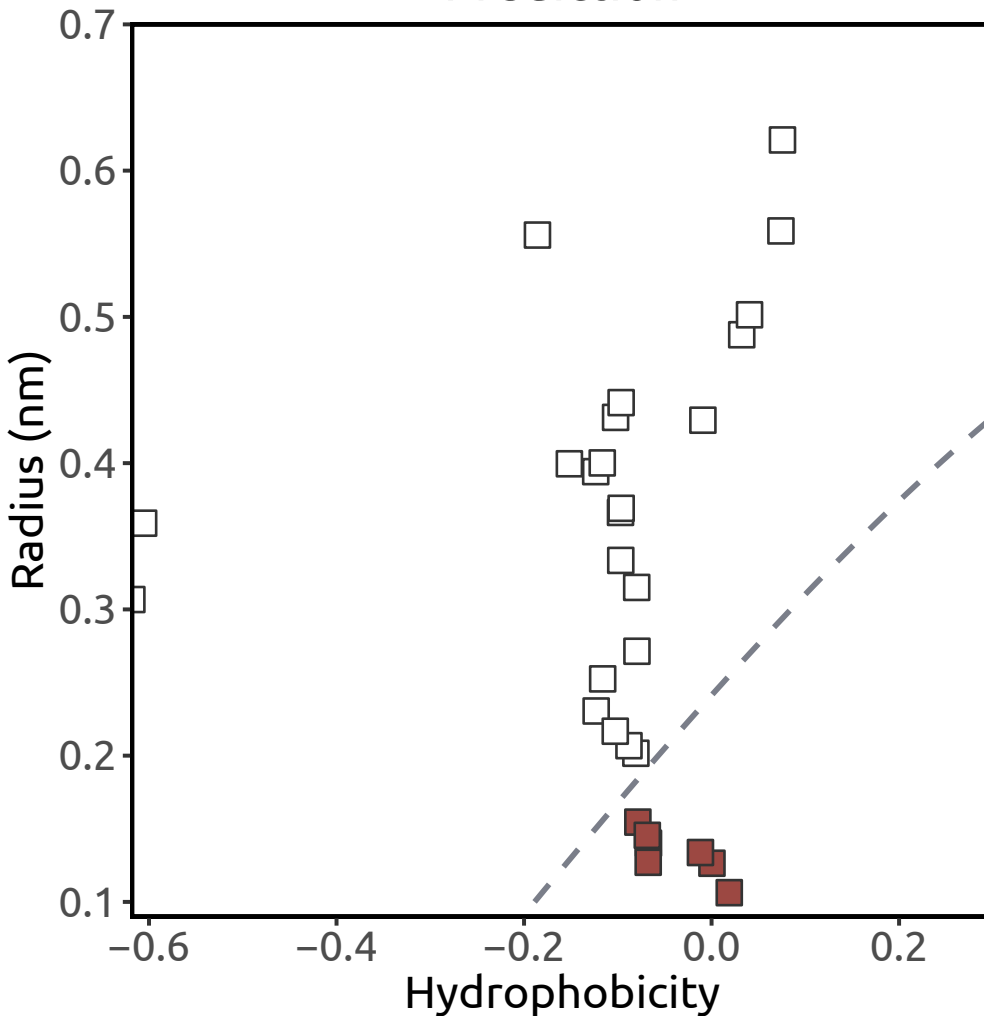
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.64 (n = 8)

Simulation result:

barrier to water

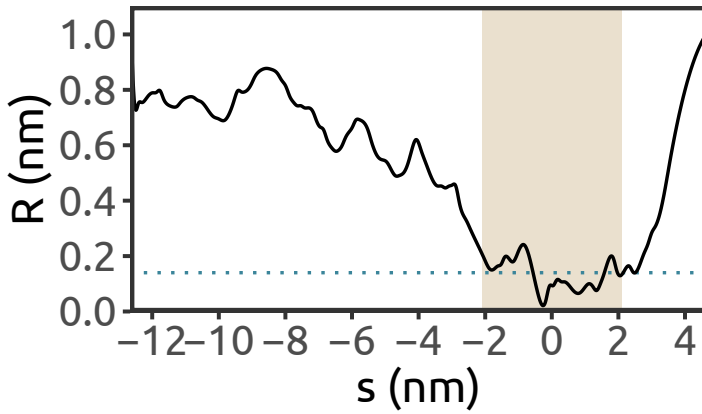
CFTR (PDB ID: 5UAR)

Danio rerio

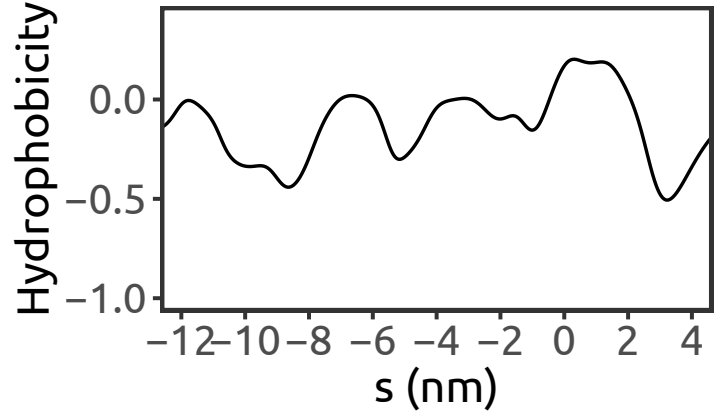
cryo-EM (3.73 Å)

Zhang & Chen, 2016

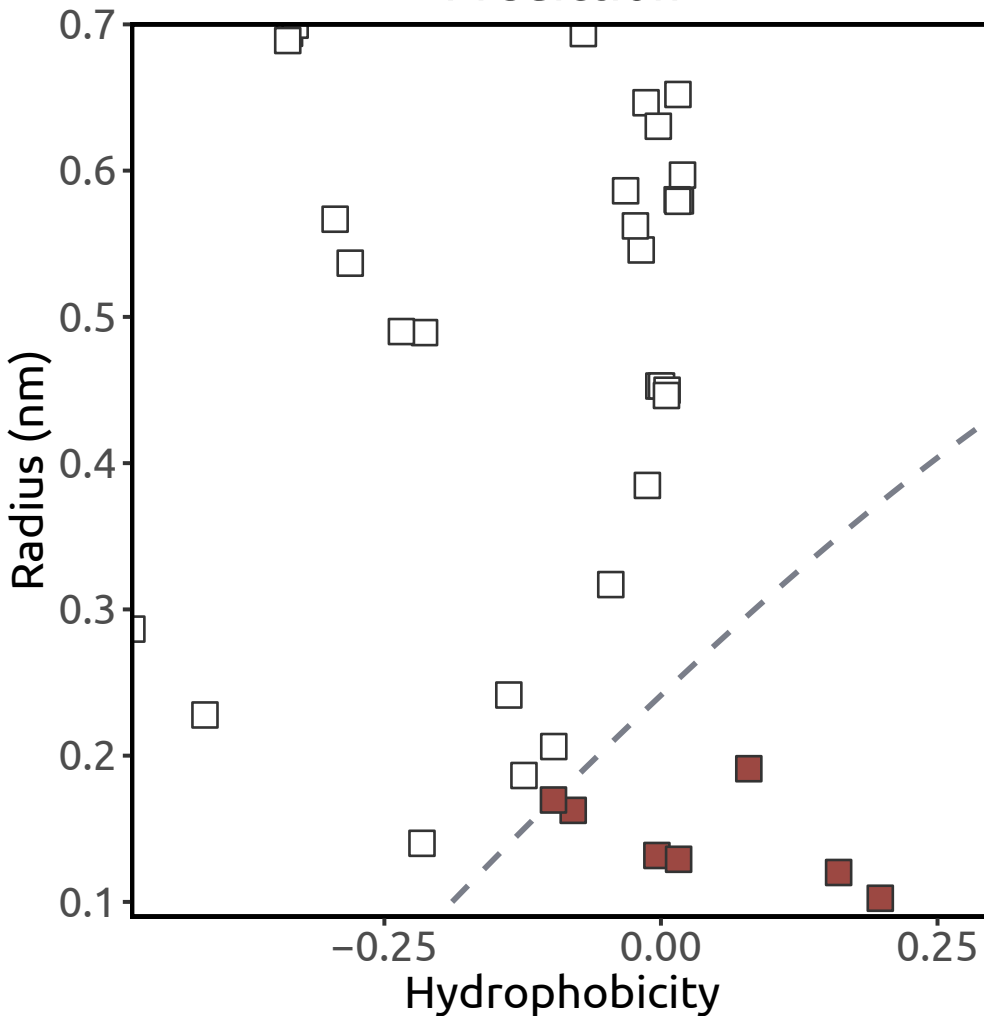
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.83 (n = 12)

Simulation result:

barrier to water

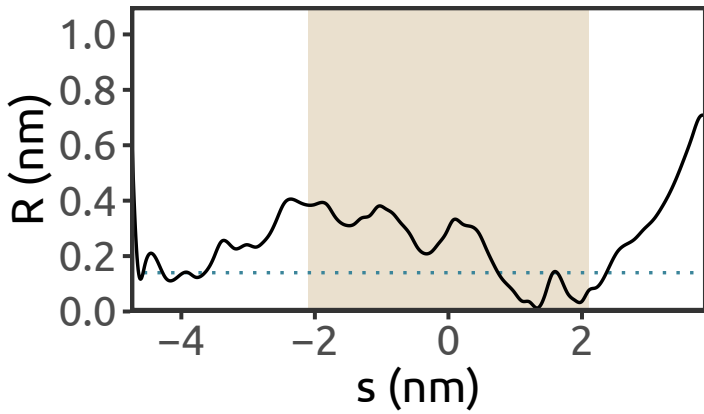
CFTR (PDB ID: 5W81)

Danio rerio

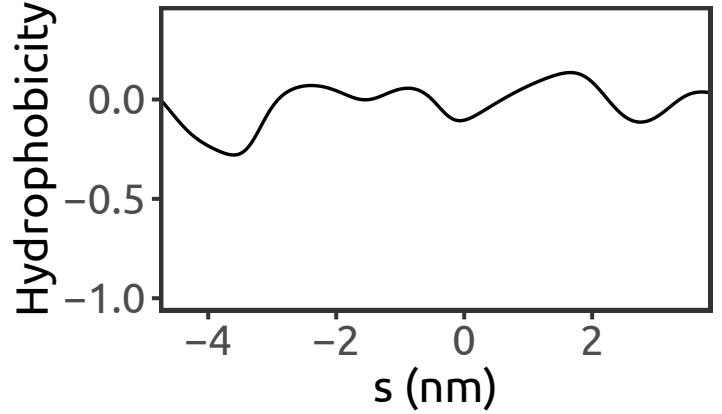
cryo-EM (3.37 Å)

Zhang et al., 2017

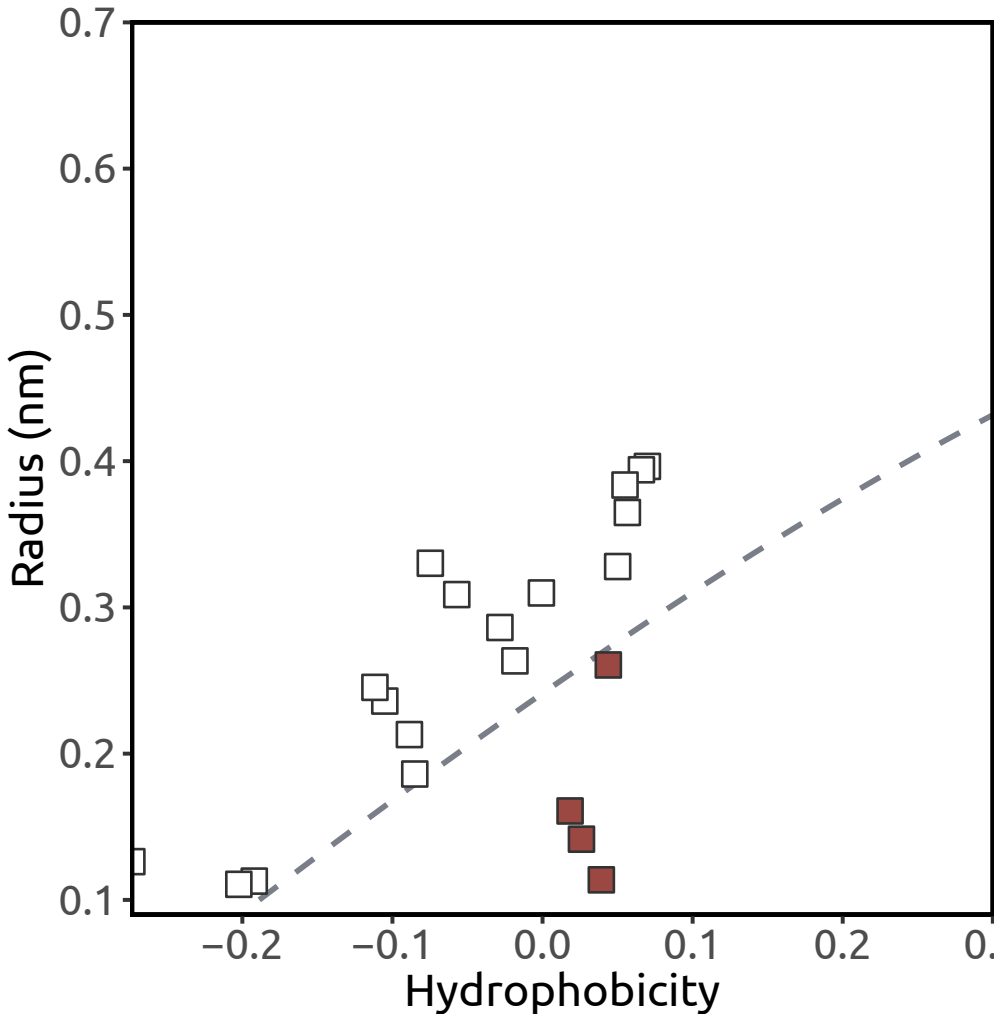
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.37 (n = 9)

Simulation result:

barrier to water

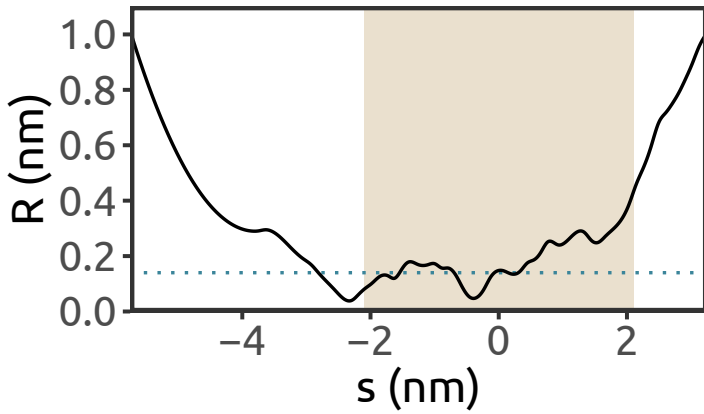
CLC (PDB ID: 5TQQ)

Bos taurus

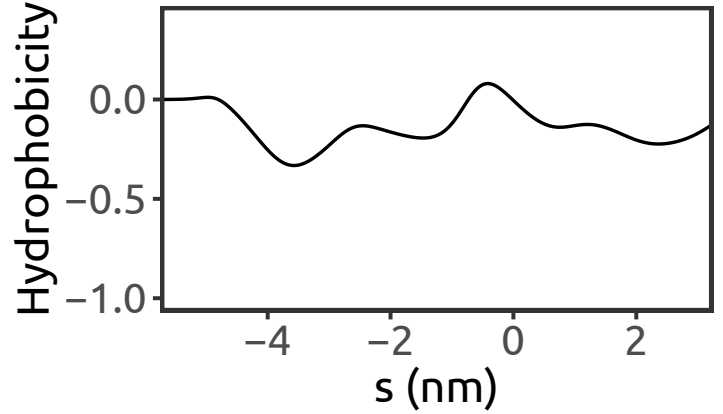
cryo-EM (3.76 Å)

Park et al., 2017

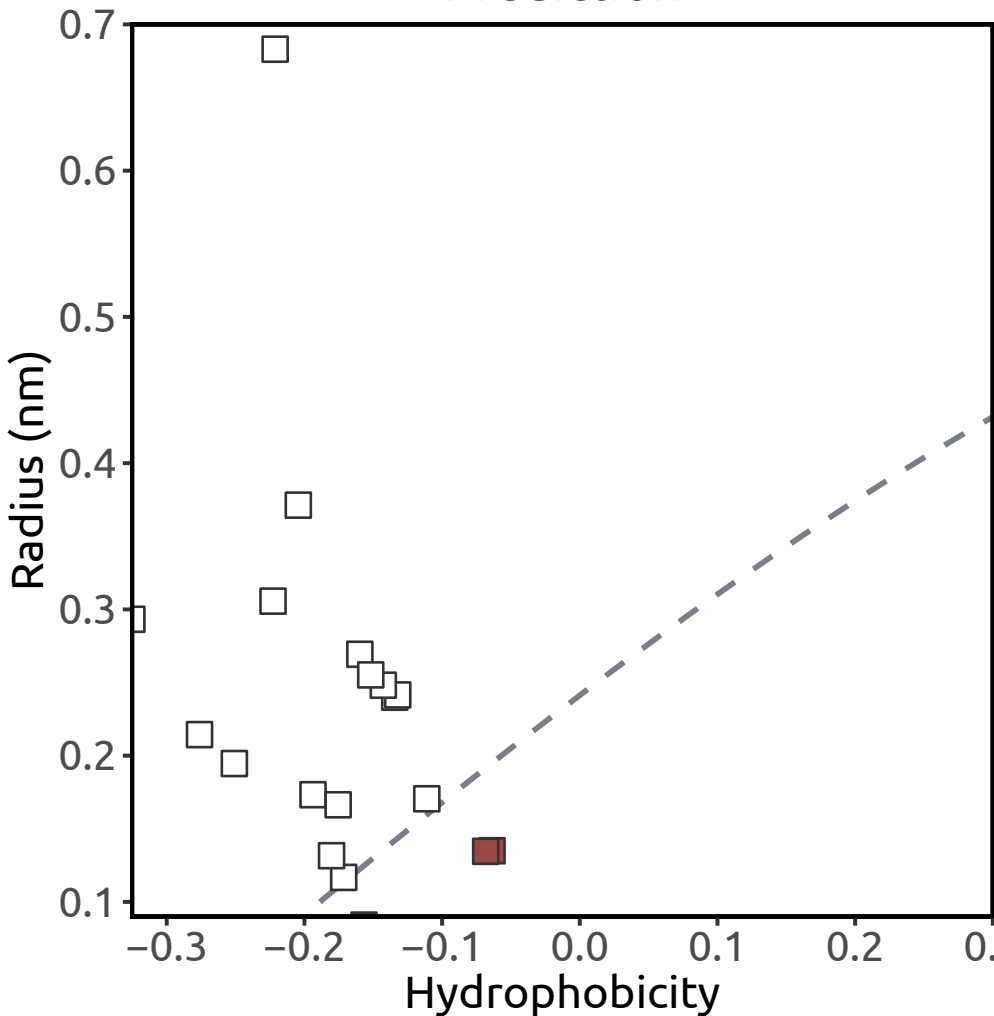
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.41 (n = 5)

Simulation result:
barrier to water

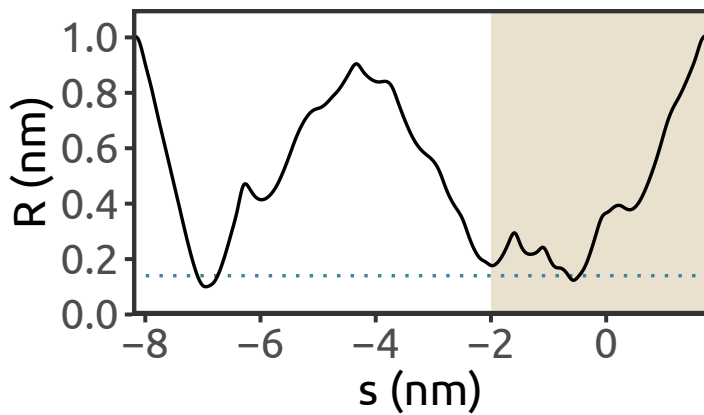
KpBest (PDB ID: 4WD8)

Klebsiella pneumoniae

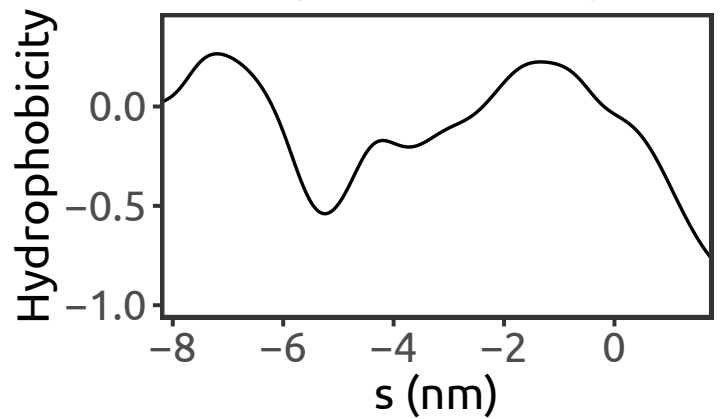
X-ray (2.3 Å)

Yang et al., 2014

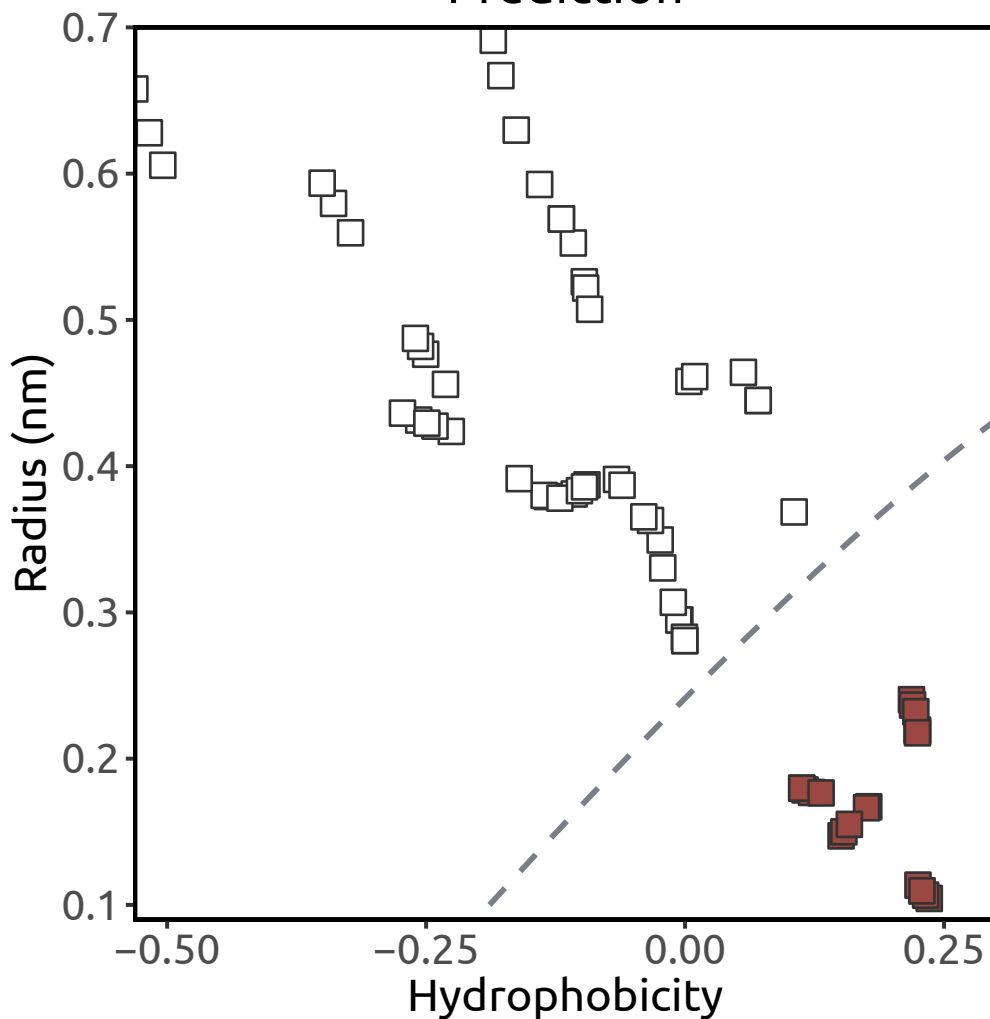
Pore radius



Hydrophobicity



Prediction



Heuristic score:
3.57 (n = 22)

Simulation result:
barrier to water

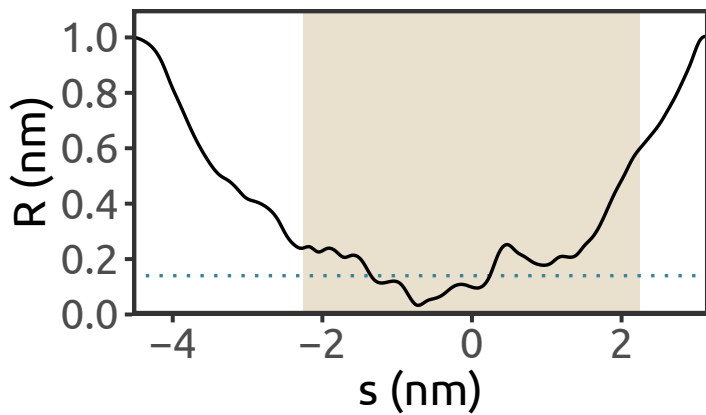
TehA (PDB ID: 3M71)

Haemophilus influenzae

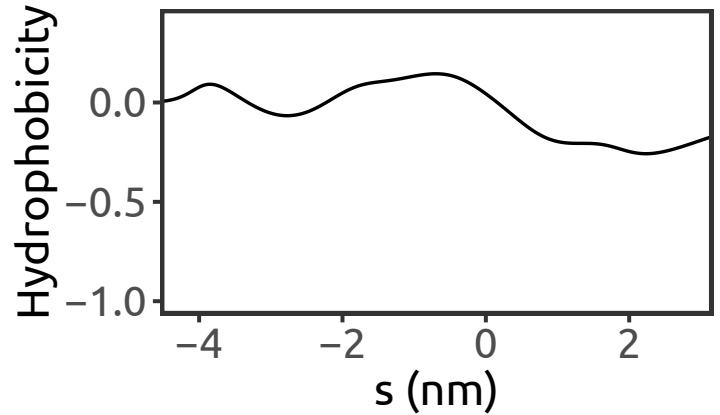
X-ray (1.2 Å)

Chen et al., 2010

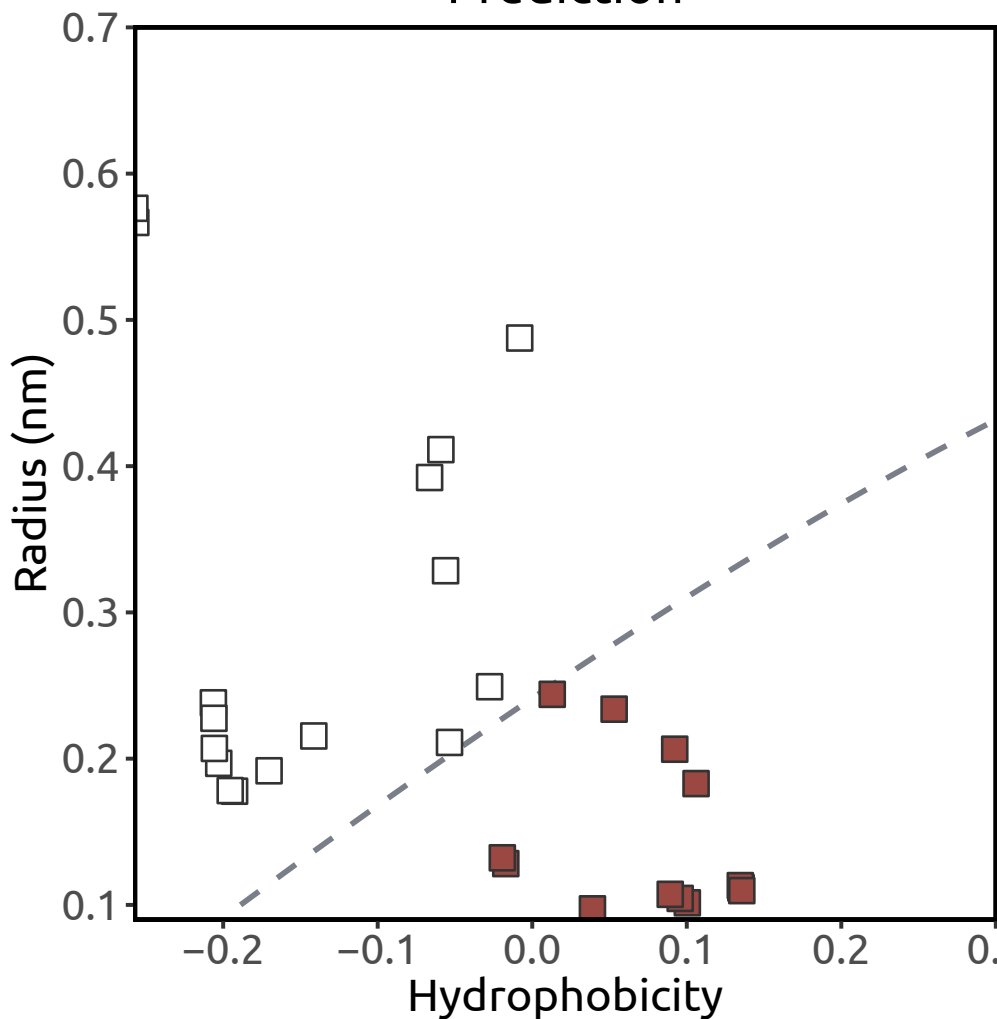
Pore radius



Hydrophobicity



Prediction



Heuristic score:
2.51 (n = 17)

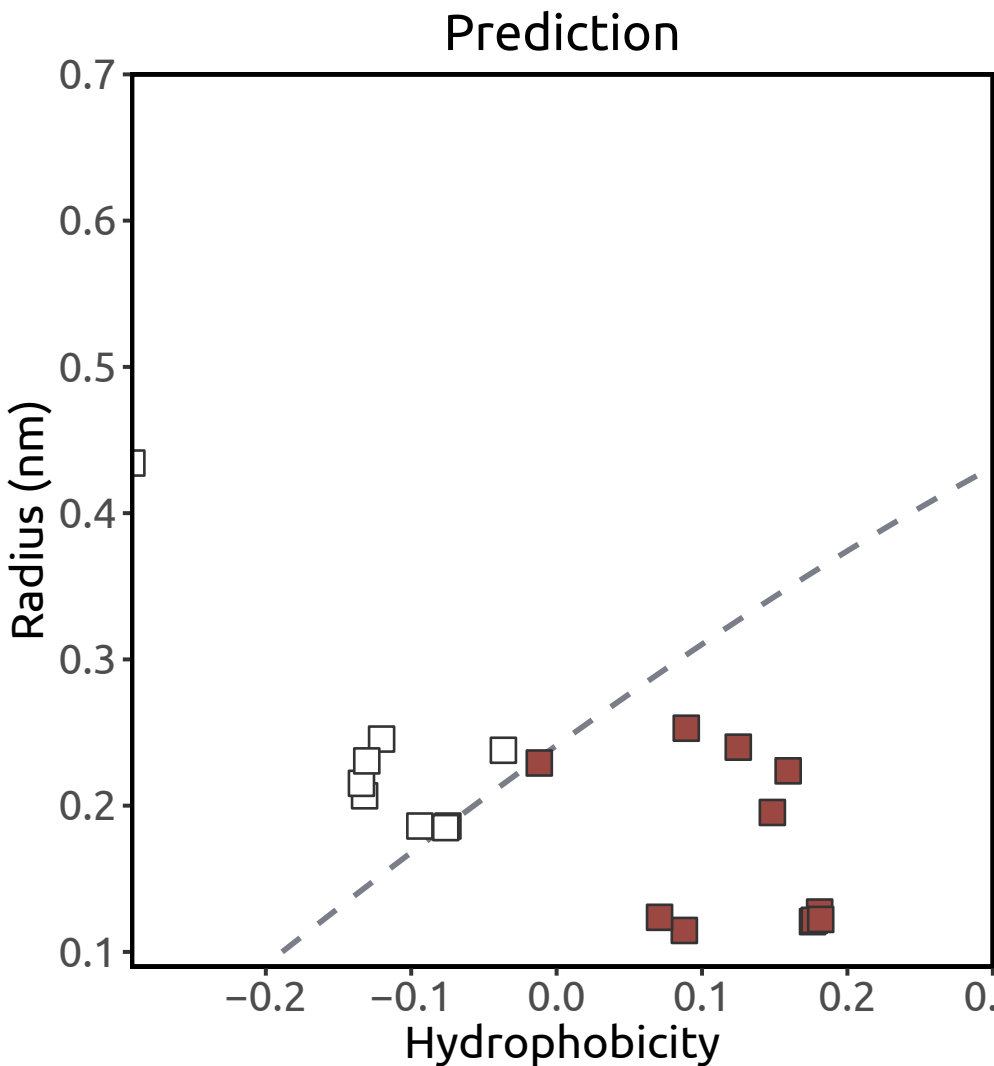
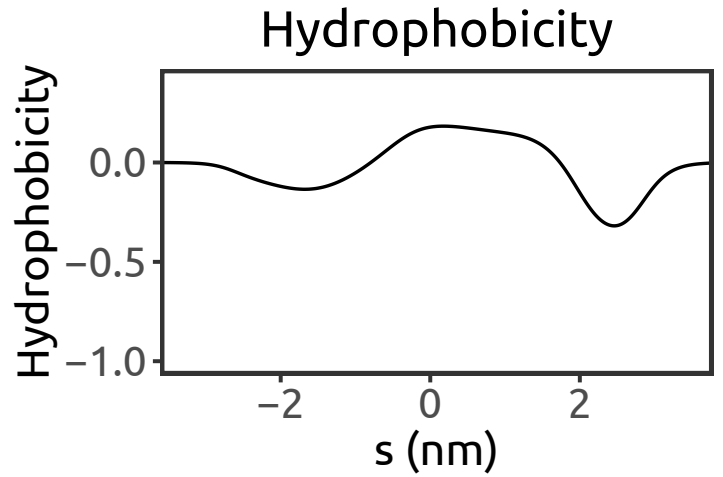
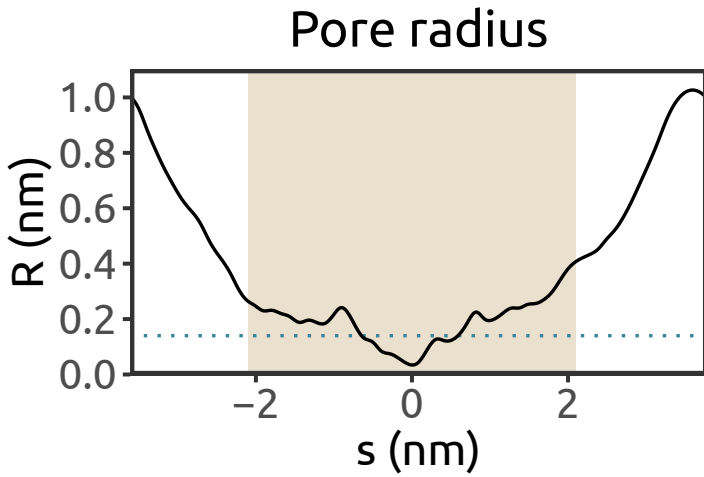
Simulation result:
barrier to water

TehA (PDB ID: 4YCR)

Haemophilus influenzae

X-ray (2.3 Å)

Axford et al., 2015



Heuristic score:
1.96 (n = 14)

Simulation result:
barrier to water

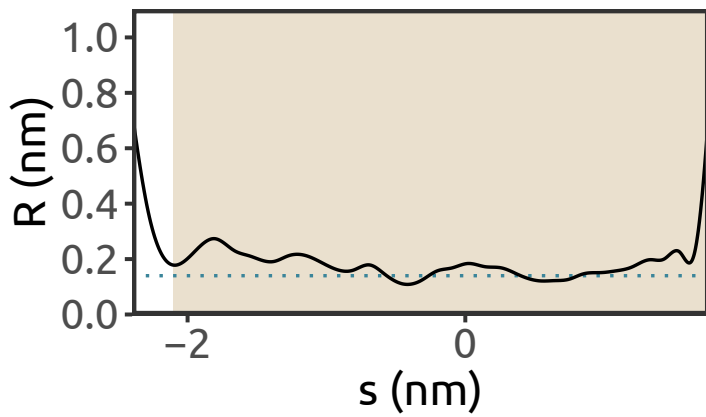
TMEM16A (PDB ID: 5OYB)

Mus musculus

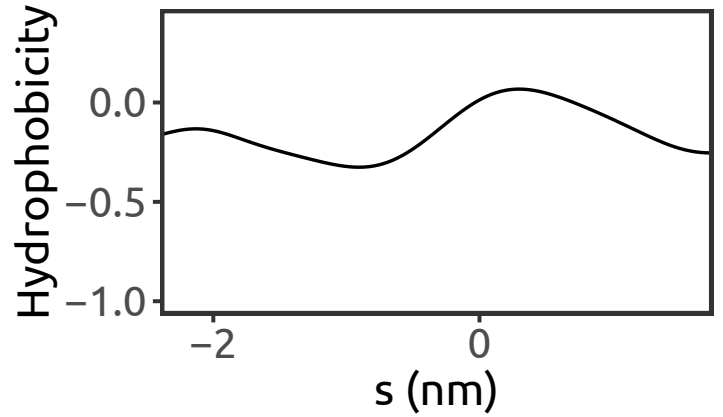
cryo-EM (3.75 Å)

Paulino et al., 2017

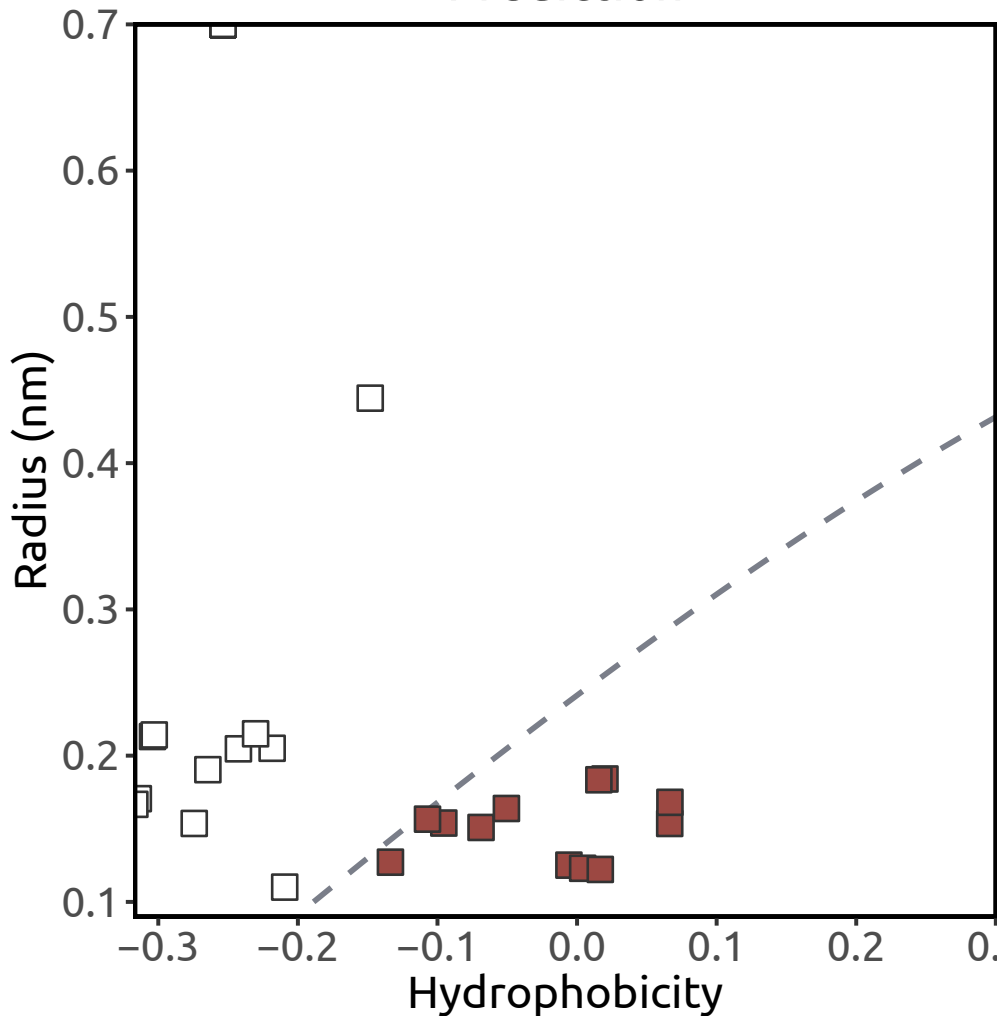
Pore radius



Hydrophobicity



Prediction



Heuristic score:

0.71 (n = 12)

Simulation result:

barrier to water

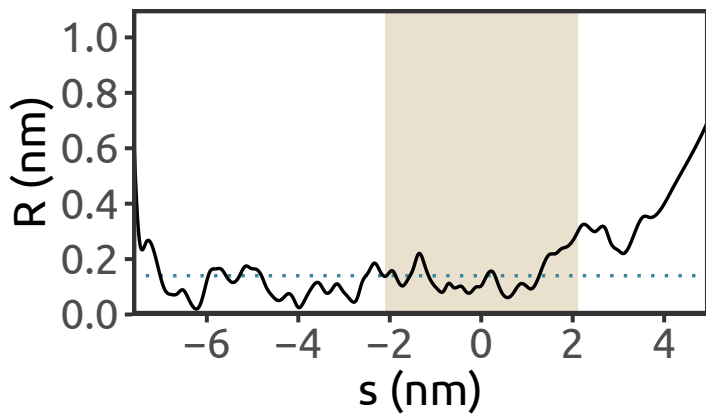
TMEM16A (PDB ID: 5OYG)

Mus musculus

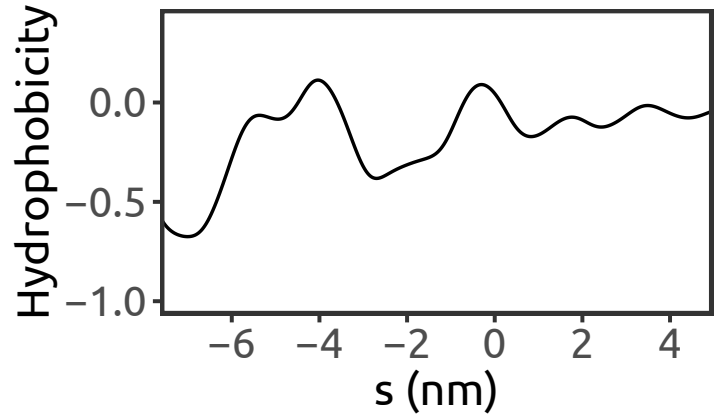
cryo-EM (4.06 Å)

Paulino et al., 2017

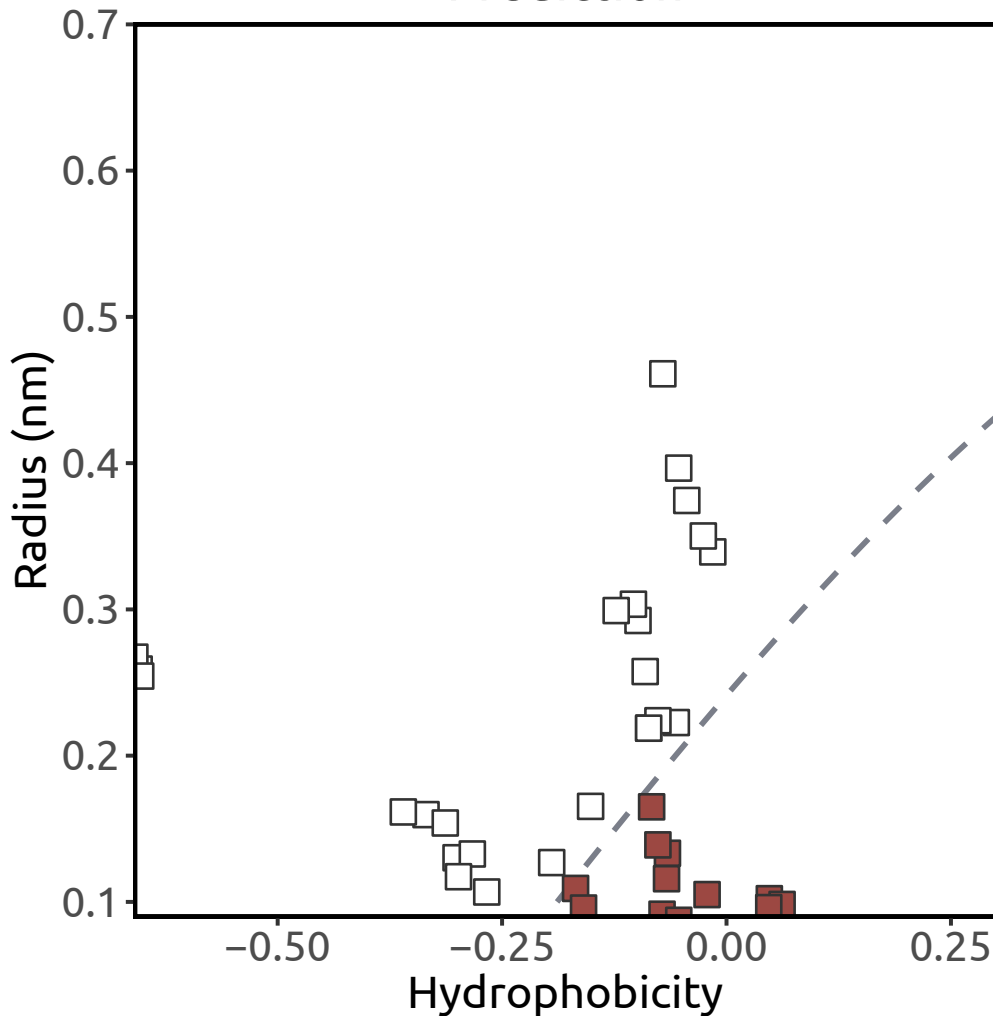
Pore radius



Hydrophobicity



Prediction



Heuristic score:

1.36 (n = 15)

Simulation result:

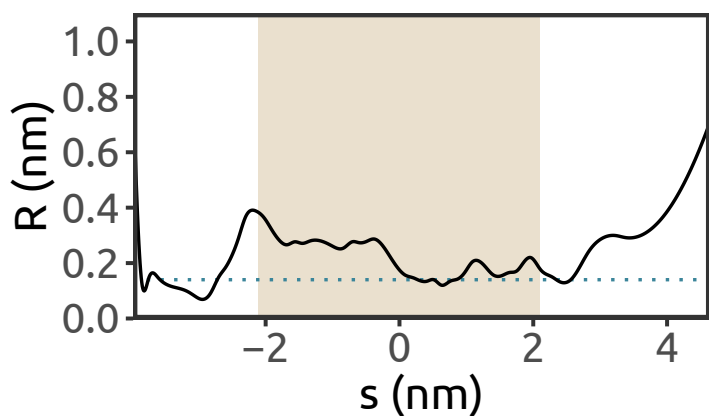
barrier to water

TMEM16A (PDB ID: 6BGI)

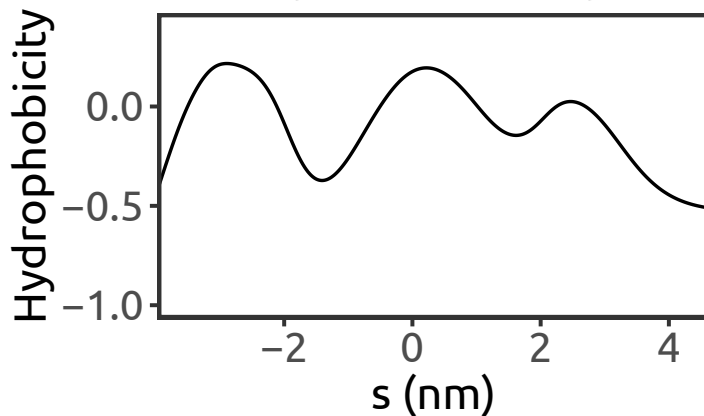
Mus musculus
cryo-EM (3.8 Å)

Dang et al., 2017

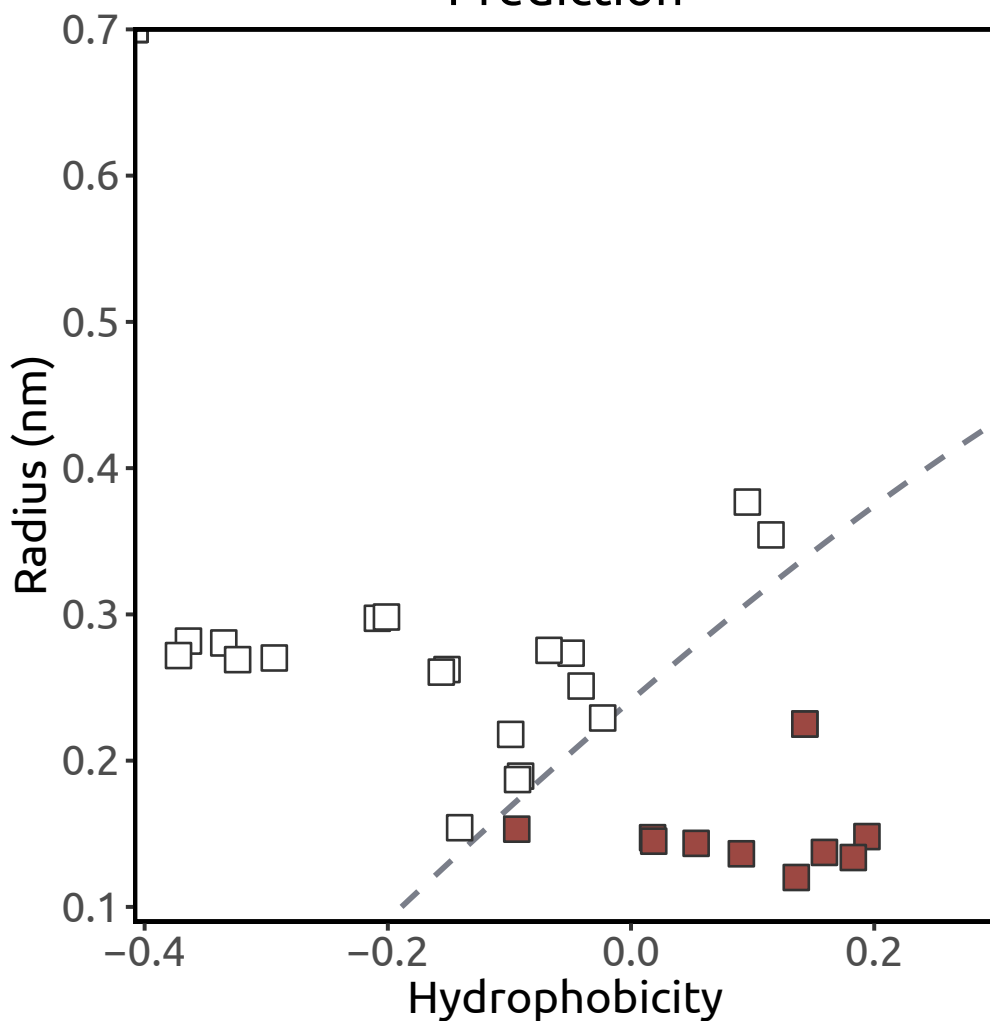
Pore radius



Hydrophobicity



Prediction



Heuristic score:
1.36 (n = 11)

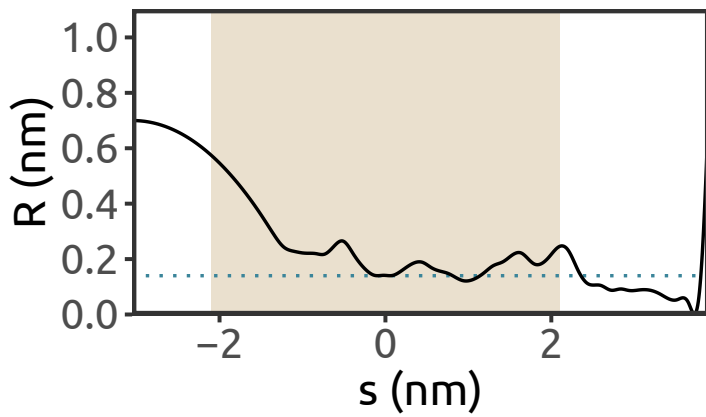
Simulation result:
barrier to water

TMEM16A (PDB ID: 6BGJ)

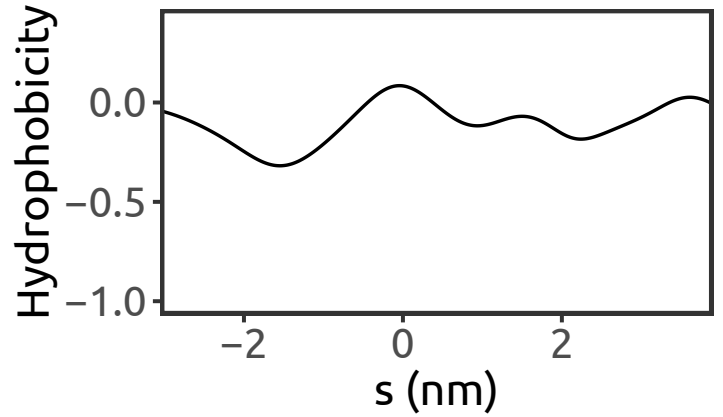
Mus musculus
cryo-EM (3.8 Å)

Dang et al., 2017

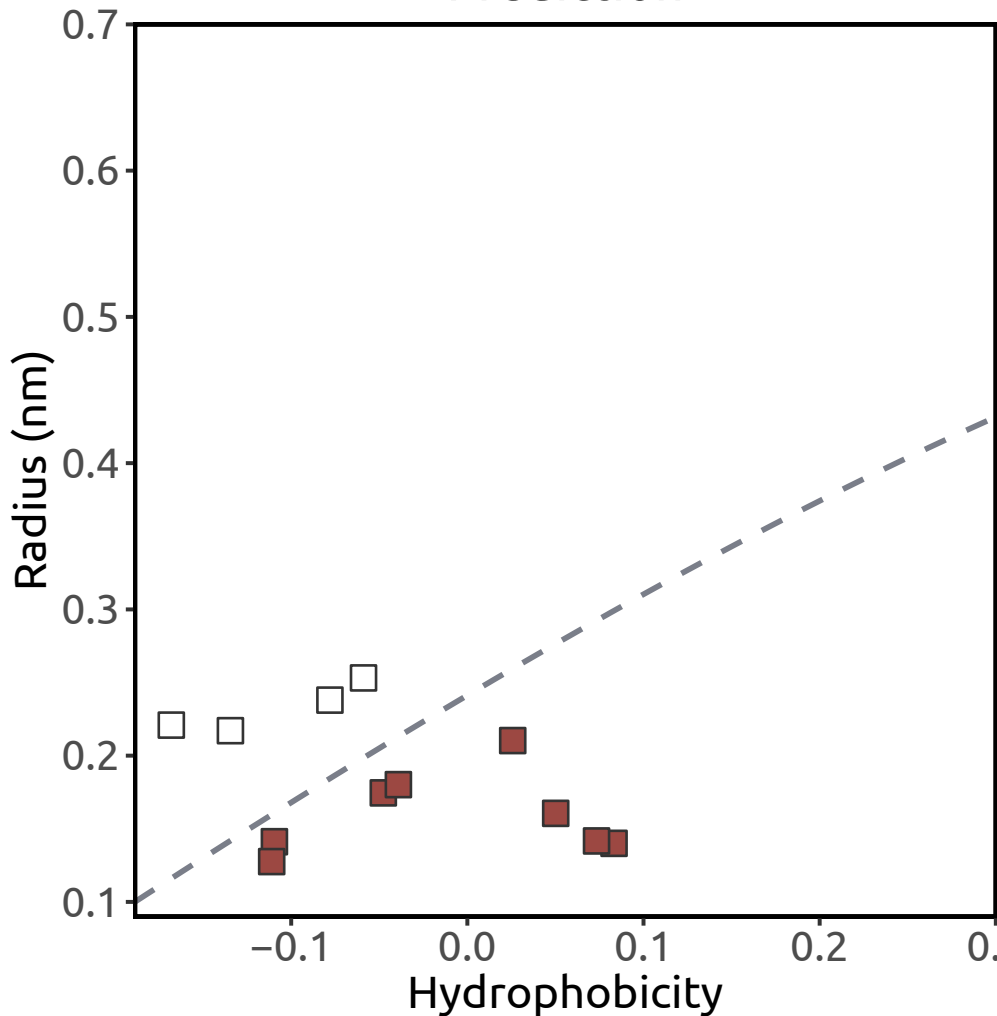
Pore radius



Hydrophobicity



Prediction



Heuristic score:
0.48 (n = 8)

Simulation result:
barrier to water