

CoEnzymes **vitamin B<sub>3</sub>** NAD<sup>+</sup>(Ox), NADH(Red). **A. Task** descriptions: for studies research of: **ADH**

**Alcohol Dehydrogenase** ChemScape MDL  **RasMol**  **MAGE**   Firefox application.

**B. Task** Lunch the molecular tutorial prepared by Aris Kaksis 2023 Riga Stradin's University look at: <http://aris.gusc.lv/ChemFiles/AlhoDeHydrogenase/NadDehydrogenase.htm> the **CPK** color scheme 1965

**1.** What N- and C-terminus amino acids have **ADH IV**? Menu Backbone, Termini Display option starting amino acid is SER.....and finishing PHE.....? What total number (see 2<sup>nd</sup> page in 1AGN.pdb) ..... and amino acids on **1JU9.pdb** crystallized structure chain.....?

**2.** What Enzyme Class (of seven Classes 1.,2.,3.,4.,5.,6.,7.) present **ADH**? .....

**3.** What particle in **ADH** transfer two reducing equivalents (**2e<sup>-</sup>**) from alcohol to **NAD<sup>+</sup>**? .....

**4.** Summary Red-Ox reaction studies (endoergic or exoergic) of **ADH** by reduced form ethanol and oxidised NAD<sup>+</sup> solutions **4.1 – 4.17 ! Absolute** potential standard values **E<sup>o</sup>** by David Harris, Kortly Shucha: at standard conditions of **absolute** scale temperature ..... K degree according to the Celsius scale 25° C .

Alcohol dehydrogenase alcohol oxidation to aldehyde (aerobic).

**4 Oks** NAD<sup>+</sup>+H<sup>+</sup>(2e<sup>-</sup>)=NADH ; -E<sup>o</sup>=**0,4095 V** **absolute inverse** standard potential David Harris.

**4 Red** CH<sub>3</sub>CH<sub>2</sub>OH+H<sub>2</sub>O=CH<sub>3</sub>HC=O+H<sub>3</sub>O<sup>+</sup>+H<sup>+</sup>(2e<sup>-</sup>); **absolute** standard potential E<sup>o</sup>=**-0.0550 V**

**4.3** Half reactions sum: NAD<sup>+</sup> + CH<sub>3</sub>CH<sub>2</sub>OH+H<sub>2</sub>O=>.....

**4.4** ΔE<sup>o</sup>=E<sup>o</sup><sub>H<sub>2</sub>O</sub>+E<sup>o</sup><sub>1</sub>=**-0.0550+0,4095=** ..... V, half reactions sum standard potential ΔE<sup>o</sup>.

**4.5** ΔG<sub>eqStandard</sub>=ΔE<sup>o</sup>•F•n=**0,3545\*2\*96485/1000=**..... kJ/mol standard free energy change.

$$1 > K_{eqstandard} = \frac{[NADH] \cdot [CH_3CHO] \cdot [H_3O^+]}{[NAD^+] \cdot [CH_3CH_2OH] \cdot [H_2O]} = e^{-\frac{\Delta G_{eqAerobi}}{R \cdot T}} = EXP(-68400/8,314/298,15) = \mathbf{1,038 \cdot 10^{-12}} = \dots\dots\dots$$

**4.6** Is favored or unfavored aerobic reaction : ..... [page 8](#) ; ;

$$\Delta G_{Hess} = \Delta G^{\circ}_{H_3O} - \Delta G^{\circ}_{CH_3CHO} - \Delta G^{\circ}_{NADH} - (\Delta G^{\circ}_{CH_3CH_2OH} + \Delta G^{\circ}_{H_2O} + \Delta G^{\circ}_{NAD^+}) = \\ = \mathbf{32,2824 + 1175,5732 - 151,549 - (75,2864 + 1059,11 - 237,191)} = \dots\dots\dots \text{ kJ/mol} \dots\dots\dots$$

Unfavored **equilibrium** constant **K<sub>eqAerobic</sub>=10<sup>-12</sup>** value shows stability in mixture.

Endothermic and endoergic ethanol **H<sub>3</sub>CCH<sub>2</sub>OH** oxidation **H<sub>3</sub>CCH=O**

**4.7** Hess free energy change positive ΔG<sub>Hess\_oxidation</sub>=..... kJ/mol , but

**4.8** minimizes ΔG<sub>min</sub>=ΔG<sub>eq</sub>=..... kJ/mol reaching aerobic equilibrium mixture:

**4.9** Aerobic oxidation with [NAD<sup>+</sup>]/[NADH]=10<sup>6</sup> homeostasis pH=7,36 is favored .

$$\Delta G_{AerobicOx} = \mathbf{68,4 + 8,3144 \cdot 298,15 \cdot \ln(1/10^6 \cdot 1/1 \cdot 10^{-(7,36)}/55,3457)/1000} = \dots\dots\dots \text{ kJ/mol} ;$$

**Inverse** symmetry: aerobic oxidation is **inverse** symmetric anaerobic reduction :

$$10^{-12} = \frac{[NADH] \cdot [CH_3CHO] \cdot [H_3O^+]}{[NAD^+] \cdot [CH_3CH_2OH] \cdot [H_2O]} = K_{eqAerobic} < 1 < K_{eqAnaerobic} = \frac{[NAD^+] \cdot [CH_3CH_2OH] \cdot [H_2O]}{[NADH] \cdot [CH_3CHO] \cdot [H_3O^+]} = 10^{12}$$

same number |ΔG<sub>Hess\_oxidation</sub>|=|.....| kJ/mol = |ΔG<sub>Hesa</sub>|=|.....| kJ/mol of opposite sign

**Inverse** exothermic and exoergic ethanal **H<sub>3</sub>CCH=O** reduction **H<sub>3</sub>CCH<sub>2</sub>OH**

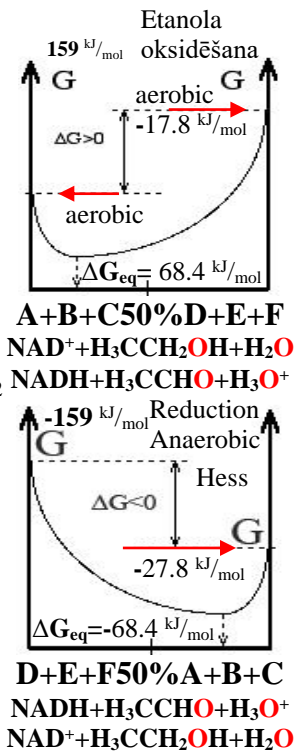
**4.10** hypoxic anaerobic ethanal reduction is **inverse** negative: ΔG<sub>Hesa</sub>=-..... kJ/mol , but

**4.11** minimized about ΔG<sub>eq</sub>=ΔE<sup>o</sup>•F•n=-0,3545 V•2 mol•96485 C/mol=-.... kJ/mol.

**4.12** Ethanal reduction about **ethanol** with anaerobic ratio [NAD<sup>+</sup>]/[NADH]=1/10 and pH=7,36 is favored, negative, exoergic free energy change i :

$$\Delta G_{anaerobic} = \mathbf{-68,4 + 8,3144 \cdot 298,15 \cdot \ln(\frac{1}{10} \cdot \frac{1}{10} \cdot \frac{55,3}{10^{-7,36}})} = \dots\dots\dots \text{ kJ/mol} ;$$

$$\Delta G_{AnaerobicRed} = \mathbf{-68,4 + 8,3144 \cdot 298,15 \cdot \ln(1/10 \cdot 1/10 \cdot 55,3457/10^{-(7,36)})/1000} = \dots\dots\dots \text{ kJ/mol} ;$$



**5.0** What **ADH IV** isoelectric point  $IEP=pH=pK_{a-vid}$  at physiologic  $pH=7,36$  ? To determine water solution  $pH$  with **ADH IV** concentration  $C=10^{-7,05339}$  M (mol/Litre)!

**Alcohol dehydrogenase ADH E.1.1.1.1. oxidoreductase**

Sequence of 386 AA amino acids for human ADH IV molecule 1AGN.pdb:

MFAEIQIQDKDRMGTAGKVIKCKAAVLWEQKQPFSEIEIEVAPPKTKEVRIKILATGICRTDDHVIKGTMVSKFPVIVGH  
 EATGIVESIGEGVTTVKPGDKVIPLFLPQCRECNACRNPDGNLCSIRSDITGRGVLDGTTTRFTCKGKPVHHFMNTSTFTE  
 YTVVDESSVAKIDDAAPPEKVKCLIGCGFSTGYGAAVKTGKVKPGSTCVVFGGLGGVGLSVIMGCKSAGASRIIGIDLNKDK  
 FEKAMAVGATECISPDKSTKPISEVLSEMTGNVGYTFEVIHLETMIDALASCHMNYGTSVVVGVPPSAKMLTYDPMLL  
 FTGRTWKGCVFGGLKSRDDVPLKLVTEFLAKKFDLDQLITHVLPFKKISEGFELLNSGQSIRTVLTF

| AA | pK <sub>aCOO-</sub> | pK <sub>aNH<sub>3</sub><sup>+</sup></sub> | pK <sub>RR</sub> | Nr | AA | pK <sub>aCOO-</sub> | pK <sub>aNH<sub>3</sub><sup>+</sup></sub> | pK <sub>RR</sub> | Nr  |
|----|---------------------|---|------------------|----|----|---------------------|---|------------------|-----|
| M  | 9,21                |   |                  | 1  | D  | 3,65                |   |                  | 59  |
| E  | 4,25                |   |                  | 2  | E  | 4,25                |   |                  | 60  |
| D  | 3,65                |   |                  | 3  | K  | 10,53               |   |                  | 61  |
| K  | 10,53               |   |                  | 4  | C  | 8,18                |   |                  | 62  |
| D  | 3,65                |   |                  | 5  | C  | 8,18                |   |                  | 63  |
| R  | 12,48               |   |                  | 6  | Y  | 10,07               |   |                  | 64  |
| K  | 10,53               |   |                  | 7  | K  | 10,53               |   |                  | 65  |
| K  | 10,53               |   |                  | 8  | K  | 10,53               |   |                  | 66  |
| C  | 8,18                |   |                  | 9  | K  | 10,53               |   |                  | 67  |
| K  | 10,53               |   |                  | 10 | C  | 8,18                |   |                  | 68  |
| E  | 4,25                |   |                  | 11 | C  | 8,18                |   |                  | 69  |
| K  | 10,53               |   |                  | 12 | K  | 10,53               |   |                  | 70  |
| E  | 4,25                |   |                  | 13 | R  | 12,48               |   |                  | 71  |
| E  | 4,25                |   |                  | 14 | D  | 3,65                |   |                  | 72  |
| E  | 4,25                |   |                  | 15 | K  | 10,53               |   |                  | 73  |
| K  | 10,53               |   |                  | 16 | D  | 3,65                |   |                  | 74  |
| K  | 10,53               |   |                  | 17 | K  | 10,53               |   |                  | 75  |
| E  | 4,25                |   |                  | 18 | E  | 4,25                |   |                  | 76  |
| R  | 12,48               |   |                  | 19 | K  | 10,53               |   |                  | 77  |
| K  | 10,53               |   |                  | 20 | E  | 4,25                |   |                  | 78  |
| C  | 8,18                |   |                  | 21 | C  | 8,18                |   |                  | 79  |
| R  | 12,48               |   |                  | 22 | K  | 10,53               |   |                  | 80  |
| D  | 3,65                |   |                  | 23 | D  | 3,65                |   |                  | 81  |
| D  | 3,65                |   |                  | 24 | K  | 10,53               |   |                  | 82  |
| H  | 6                   |   |                  | 25 | E  | 4,25                |   |                  | 83  |
| K  | 10,53               |   |                  | 26 | E  | 4,25                |   |                  | 84  |
| K  | 10,53               |   |                  | 27 | Y  | 10,07               |   |                  | 85  |
| H  | 6                   |   |                  | 28 | E  | 4,25                |   |                  | 86  |
| E  | 4,25                |   |                  | 29 | H  | 6                   |   |                  | 87  |
| E  | 4,25                |   |                  | 30 | E  | 4,25                |   |                  | 88  |
| E  | 4,25                |   |                  | 31 | D  | 3,65                |   |                  | 89  |
| K  | 10,53               |   |                  | 32 | C  | 8,18                |   |                  | 90  |
| D  | 3,65                |   |                  | 33 | H  | 6                   |   |                  | 91  |
| K  | 10,53               |   |                  | 34 | Y  | 10,07               |   |                  | 92  |
| C  | 8,18                |   |                  | 35 | K  | 10,53               |   |                  | 93  |
| R  | 12,48               |   |                  | 36 | Y  | 10,07               |   |                  | 94  |
| E  | 4,25                |   |                  | 37 | D  | 3,65                |   |                  | 95  |
| C  | 8,18                |   |                  | 38 | R  | 12,48               |   |                  | 96  |
| C  | 8,18                |   |                  | 39 | K  | 10,53               |   |                  | 97  |
| R  | 12,48               |   |                  | 40 | C  | 8,18                |   |                  | 98  |
| D  | 3,65                |   |                  | 41 | K  | 10,53               |   |                  | 99  |
| C  | 8,18                |   |                  | 42 | R  | 12,48               |   |                  | 100 |
| R  | 12,48               |   |                  | 43 | D  | 3,65                |   |                  | 101 |
| D  | 3,65                |   |                  | 44 | D  | 3,65                |   |                  | 102 |
| R  | 12,48               |   |                  | 45 | K  | 10,53               |   |                  | 103 |
| D  | 3,65                |   |                  | 46 | E  | 4,25                |   |                  | 104 |
| R  | 12,48               |   |                  | 47 | K  | 10,53               |   |                  | 105 |
| C  | 8,18                |   |                  | 48 | K  | 10,53               |   |                  | 106 |
| K  | 10,53               |   |                  | 49 | D  | 3,65                |   |                  | 107 |
| K  | 10,53               |   |                  | 50 | D  | 3,65                |   |                  | 108 |
| H  | 6                   |   |                  | 51 | H  | 6                   |   |                  | 109 |
| H  | 6                   |   |                  | 52 | K  | 10,53               |   |                  | 110 |
| E  | 4,25                |   |                  | 53 | K  | 10,53               |   |                  | 111 |
| Y  | 10,07               |   |                  | 54 | E  | 4,25                |   |                  | 112 |
| D  | 3,65                |   |                  | 55 | E  | 4,25                |   |                  | 113 |
| E  | 4,25                |   |                  | 56 | R  | 12,48               |   |                  | 114 |
| K  | 10,53               |   |                  | 57 | F  | 1,83                |   |                  | 115 |
| D  | 3,65                |   |                  | 58 |    |                     |   |                  |     |

115 of 386 amino acids active values pKa

Sum = 881,66.....

=  $\Sigma pK_{aRside\ group} + pK_{aNterminal} + pK_{aCterminal} =$

$pK_{amean} = (\Sigma pK_{aRside\ group} + pK_{aNterminal} + pK_{aCterminal}) / NpKa$

$IEP = pK_{amean} = 881,66 / 115 = 7.6666.....$

Calculation tasks for human ADH IV molecule 1AGN.pdb

Protolytic constant, isoelectric point IEP=  $pK_{a\text{mean}}$  calculate of side chains  $\Sigma pK_{a\text{Rside group}}$ .  $pK_{a\text{Nterminal}} \text{NH}_3$

and  $pK_{a\text{Cterminal}} \text{COO}^-$  constants sum divide with number of acid groups  $NpK_a$ :

$$\text{IEP} = pK_{a\text{mean}} = (\Sigma pK_{a\text{Rside group}} + pK_{a\text{Nterminal}} + pK_{a\text{Cterminal}}) / NpK_a$$

1 Acid groups number in sum  $NpK_a = 113 + 2 + \dots = \dots$

386 amino acids of them protolytic constants  $pK_a$  for side groups  $113 + 2$  terminus N and C,

N-terminal methionine M  $pK_{a\text{Nterminal}} = 9.21$  and C-terminal phenylalanine F  $pK_{a\text{Cterminal}} = 1.83$

Sum are calculating as  $\Sigma pK_{a\text{Rside group}} + pK_{a\text{Nterminal}} + pK_{a\text{Cterminal}} = \dots$

2 Average acid group constant  $pK_{a\text{mean}} = \text{IEP}$  **ISOELEKTRIC POINT**

$$\text{IEP} = pK_{a\text{mean}} = 881,66 / 115 = \dots$$

At pH value of amino acid and protein on isoelectric point  $\text{pH} = \text{IEP}$  total charge is zero „0”

0 — plus (+) acidic — zero charge „0”  $\text{IEP} = \text{pH}$  — minus (-) basic — 14 pH scale

$-\text{COOH}$  &  $-\text{NH}_3^+$  positive charge .....  $-\text{COO}^-$  &  $-\text{NH}_2$  ..... charge is negative  $-\text{COO}^-$  &  $-\text{NH}_2$

Underline and determine existing: positive (+) or negative (-) or zero !

3 Determine ADH IV molecule charge sign (+), zero „0” or (-) at physiologic  $\text{pH} = 7.36$

Underline existing:

$-\text{COOH}$  &  $-\text{NH}_3^+$  positive (+) charge .....  $\text{pH} = 7.36 < \text{IEP} = 7.67$  ..... charge negative (-)  $-\text{COO}^-$  &  $-\text{NH}_2$ .

4 Determine ADH IV molecule charge sign (+), zero „0” or (-) at **electrophoresis**  $\text{pH} = 8.8$

Underline existing:

$-\text{COOH}$  &  $-\text{NH}_3^+$  positive (+) charge .....  $\text{IEP} = 7.67 < \text{pH} = 8.8$  ..... charge negative (-)  $-\text{COO}^-$  &  $-\text{NH}_2$ .

5 Calculate ADH IV solution pH at concentration  $C = 10^{-7,05339} \text{ M}$  ( mol / Litre)

by *Ostwald dilution law* concentration M in logarithm:

$$\text{pH} = \frac{pK_{a\text{mean}} - \log C}{2} = \frac{7.6666087 - \log 10^{-7,0533913}}{2} = \frac{7.6666087 + 7,0533913}{2} = 14,72 / 2 = \dots$$

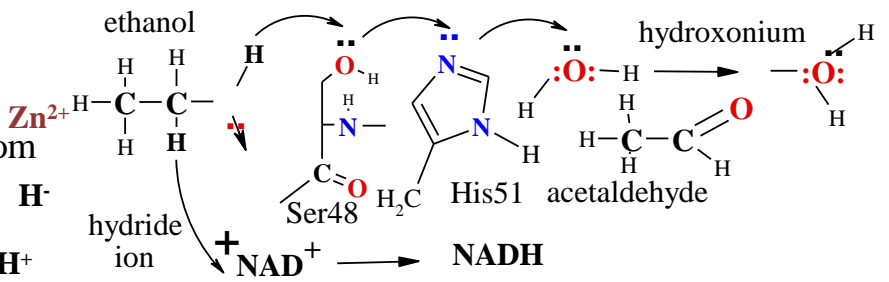
7,36 Attractor ADH IV concentration is  $C = \dots \text{ M}$ .

5. . Place catalytic  $\text{Zn}^{2+}$  ion, ethanol oxygen atom O coordinate with donor acceptor bond and four jumping dissociated proton  $\text{H}^+$  pathway from alcohol group  $-\text{CH}_2-\text{O}-\text{H}$  to Ser48 to His51 and resulting bound proton  $\text{H}^+$  to water molecule  $\text{H}_2\text{O}$  forming hydronium ion  $\text{H}_3\text{O}^+$ .

**ADH during alcohol oxidation in water medium ,**

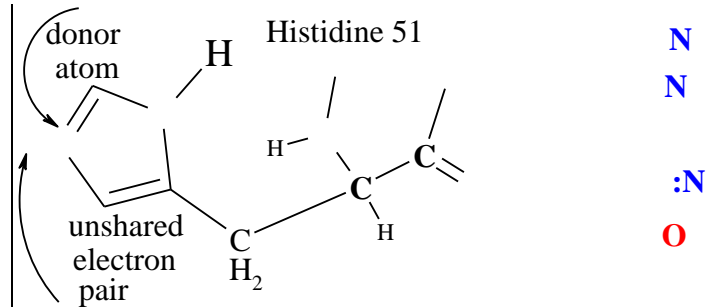
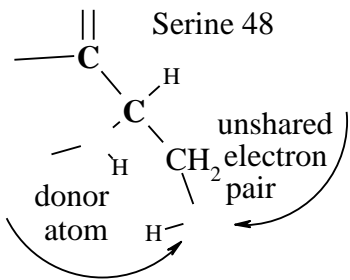
Place hydride ion  $H^-$  in to tunneling pathway from alcohol group carbon atom- $CH_2$ - to  $NAD^+$  cyclic carbon atom

$-HC^+-$  producing NADH



that  $H_2O$  water molecule forming hydroxonium ion  $H_3O^+$  and aldehyde

6. Place in **Ser-48, His-51** structures  $O$ ,  $N$  atoms and electrons pair donor atoms  $O:$ ,  $:N$ !



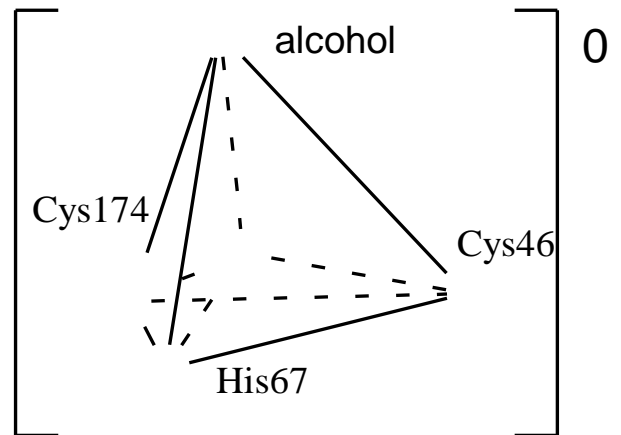
7. Place  $O$ ,  $Zn^{2+}$ ,  $S$ ,  $N$  atoms and charge zero 0 of complex  $[Zn^{2+}(S-Cys)_2(O-spirts)(NHis)]^0$

for tetragonal geometry,

like trigonaal pyramid!

1HLD.pdb  $Zn^{2+}$  ion coordinates:

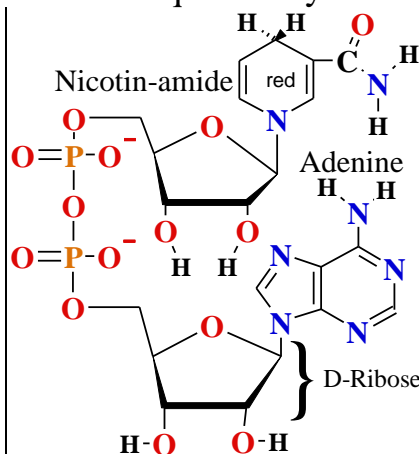
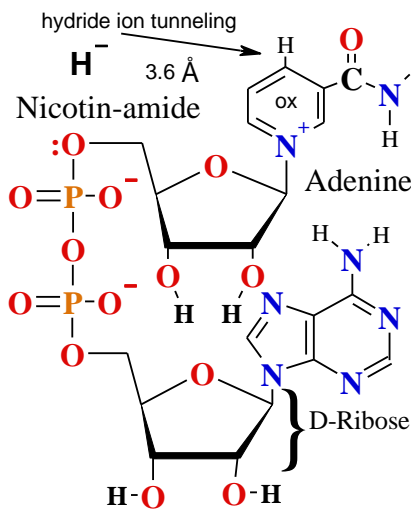
Cys46-Cys174-His67- $O$  alcohol



8. What vitamin-cofactor oxidizes alcohols in **ADH dimer** quaternary structure?.....

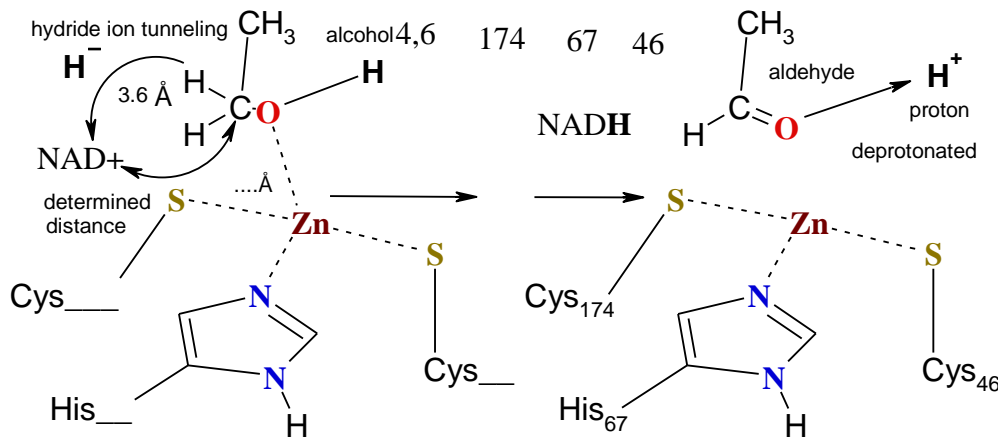
9. Oxidised  $NAD^+$  hydride tunneling **nicotine adenine dinucleotide:**

Nicotin-amide, Adenine, two riboses, two phosphates with anhydride bond between phosphates



10. **NADH hydride nicotine adenine dinucleotide** reduced form: Nicotin-amide, Adenine, two riboses, two phosphates with anhydride bond between phosphates

ADH 1HLD.pdb  $Zn^{2+}$  coordinates Cys46-Cys174-His67- $O$  : Tunnel distance 3,6 Å for hydride



ion  $H^-$  to  $NAD^+$  nicotine  
amide positive charged cycle  
carbon atom  $-CH-$ .  
Measure distance ..... Å from  
alcohol  $-CH_2-$  carbon atom  
to  $NAD^+$  aromatic cycle  $-CH-$   
in 1HLD.pdb molecule.  
With right button click in  
menu choose „Distance”

from „Select Mouse Click Action” measure distance from alcohol carbon atom  $-CH_2-$  ..... Å  
to  $NAD^+$  nicotine amide cyclic carbon atom  $-CH-$ !

11. Place amino acid numbers for coordination sphere and measure distance in angstroms units.

12. Secondary structures in ADH are.....helixes and..... sheets.

13. Count **alpha-helixes** on ADH polypeptide molecule? ..... **alpha-helixes**.....

14. Count **beta strands - sheets** in ADH molecule?.....**beta strands in-sheet**

..... **beta strands in-sheet** and .....**beta strand**.....

15. Count quaternary 4° structure components of 3° subunits in ADH molecule **1JU9zn.pdb**  
and **1HLDznNAD.pdb**? identical ADH molecules ....., each bind .....

each in domain binds **substrate alcohol group like** .....,.....

16. What physiological functions in human body have ADH against ethanol?.....

.... remove two hydrogen atoms and so oxidize .....

18. What toxic physiological functions in human body have ethanol molecules at long time  
abused? CSDD Car transport certification limited concentration in blood 0.5 promilles and  
more? a) slow down the transport through membrane aquaporin-channels of ...+...

b).... long time abused in body leads in tissues to .....and .....

c).... ethanol to prevent formation and compete with .....

19. What toxic functions in human body have ADH against methanol?

to poison human body with .....

remove two hydrogen atoms of alcohol so to oxidize .....

20. Complete the oxidation reaction for methanol:  $H_3C-OH+NAD^+$  in water.

$H_3C-OH+NAD^++H_2O(His51)\rightleftharpoons$ .....

Methanol B<sub>3</sub> vitamin

Formaldehyde B<sub>3</sub> vitamin reduced

21. How compete ethanol with methanol? What ist the anti dot against methanol misuse in  
human body? high ethanol concentration oppress .....

....methanol oxidation silencing lets throw aquaporins .....

22. To call six crystalline shapes for ADH subunits designation by Greek alphabet letters!

1. alpha ....., 2. beta ....., 3. gamma....., 4. pee....., 5. chi....., 6. sigma.....

23. What kind human alcohol dehydrogenase crystallization failed? .....

To depict what kind of human alcohol dehydrogenases seven types - proteins identified in organism from data bank of Uni-Prot KB files:

1. ....\_HUMAN, subunits designated .....
2. ....\_HUMAN, subunits designated .....
3. ....\_HUMAN, subunits designated .....
4. ....\_HUMAN, subunits designated .....
5. ....\_HUMAN, subunits designated .....
6. ....\_HUMAN, subunits designated .....
7. ....\_HUMAN, .....

<http://aris.gusc.lv/ChemFiles/AlhoDeHydrogenase/4DXH5VJ5hOhBioChem1718/5VJ5hOhBioChem17.pdf>

| The Class                 | System | Protein gene | Uni-Prot KB | GeneGene New Old | <b>Table 1:</b> Nomenclature for Human Alcohol Dehydrogenase <b>Abstract Background</b>  |
|---------------------------|--------|--------------|-------------|------------------|--|
| Class I <sub>1HSO</sub>   |        | ADH1A        | .           | ADH1 ADH1A       | All known attempts to isolate and characterize mammalian class V alcohol dehydrogenase (class V ADH), a member of the large ADH protein family, at the protein level |
| Class I <sub>1DEH</sub>   |        | ADH1B        | .           | ADH2 ADH1B       |  |
| Class I <sub>1HT0</sub>   |        | ADH1C        | .           | ADH3 ADH1C       |  |
| Class II                  |        | ADH2         | .           | ADH4 ADH4        |  |
| Class III <sup>1MP0</sup> |        | ADH3         | .           | ADH5 ADH5        |  |
| Class IV <sup>1AGN</sup>  |        | ADH4         | .           | ADH7 ADH7        |  |
| Class V                   |        | ADH5         | .           | ADH6 ADH6        |  |

have failed. This indicates that the class V ADH according Uni-Prot KB ADH6\_HUMAN protein is not stable in a non-cellular environment, which is in contrast to all other human ADH enzymes. In this report we present evidence, supported with results from computational analyses performed in combination with earlier in vitro studies, why this ADH behaves in an atypical way.

[Arch Biochem Biophys.](#) 2018;653:97-106. **4DXHa**

*Biochemistry*, 2017, 56 (28), pp 3632-3646.

[5ENV,8ADH,1QLH,4DWV,1N92,1N8K,1P1R,4DXH,1N92,1N8K,1LDE,1LDY,1MGO,5VKR,1HEU,2JHF,1HET,2JHG,1H2B,1MAO,1PL6,1PL6,1YKF,1YE3,4XD2,5VJ5,5VJG,5VKR,5VL0,5VN1](#)..., 6