Vol. 16, No. 1 (2017) 33-39 Revista Mexicana de Ingeniería **Q**uímica



ENZYMATIC REDUCTION BY ALCOHOL DEHYDROGENASE TA1316 FROM

Thermoplasma acidophilum

REDUCCIÓN ENZIMÁTICA POR ALCOHOL DESHIDROGENASA TA1316 PROVENIENTE DE Thermoplasma acidophilum

M. Guzmán-Rondríguez, L. Santos*

División de Biología Molecular, Instituto Potosino de Investigación Científica y Tecnológica, A.C. (IPICYT), Camino a la Presa San José 2055, Lomas 4a Sección, CP 78216, San Luis Potosí, México. Universidad Politécnica de Pachuca, Carr. Pachuca-Cd. Sahagún Km. 20, Zempoala, Hidalgo. México.

Received January 14, 2016; Accepted November 25, 2016

Abstract

In this work we present the NADH-dependent Ta1316 alcohol dehydrogenese (ADH) characterization for its reducing reaction in the presence of aldehydes, ketones and keto-esters. In the presence of 2 mM acetaldehyde, Ta1316 ADH showed a remarkable thermal activity, displaying activity at temperatures up tp 90°C and low pH with a requirement of Z^{+2} to reach its maximum activity. In contrast to other characterized ADHs, Ta1316 ADH reduces methyl pyruvate, an alpha-ketoester as its preferred substrate. We conclude that Ta1316 ADH's principalfunction is the reduction of substrates and is potentially an enzyme that can be used in biotechnology, as erll as in industrial applications.

Keywords: alcohol dehydrogenase, Ta1316, Thermoplasma acidophilum, reduction reaction, characterization.

Resumen

En este trabajo presentamos la caracterización de la actividad enzimática reductora de la enzima alcohol deshidrogenasa (ADH) Ta1316 dependiente de NADH en presencia de aldehídos, cetonas y cetoésteres. La ADH Ta1316 mostró estabilidad térmica notable con actividad máxima a temperatura de 90°C y pH acídico en presencia del cofactor Zn⁺² y acetaldehído 2 mM. En contraste con otras ADHs caracterizadas, Ta1316 es capaz de reducir metil-piruvato, un alfa-cetoéster como su sustrato preferido. Concluimos que la función principal de la Ta1316 ADH es la reducción de sustratos y es una enzima que podría ser utilizada potencialmente en aplicaciones biotecnológicas e industriales.

Palabras clave: alcohol deshidrogenasa, Ta1316, Thermoplasma acidophilum, reacción reductora, caracterización.

1 Introduction

Alcohol dehydrogenases (ADHs) are part of the ubiquitous oxidoreductase family within the three domains of life. This cofactor-dependent enzyme catalyzes the oxidation of alcohols in the presence of NAD+ or NADP+ (Reid & Fewson, 1994). ADHs are involved in the interconversion between alcohols, aldehydes and ketones that are key in biotechnological processes such as the production of optical active alcohols for fragrances, pharmaceuticals and agrochemicals. These enzymes are also useful in bioelectrocatalysis for the production of biosensors, biofuels (Eichler, 2001) (Kroutil *et al.*, 2004) (Yakushi & Matsushita, 2010) (Reyes-Valadez, 2016),

bioreduction (Yang et al., 2012), optically active compounds (Nie et al., 2007; Nie et al., 2007) or fermented spirits (De los Rios-Deras et al., 2015).

Hence, the demand in industry for active and stable ADHs; able to tolerate elevated temperatures, organic solvents or distinct pHs has increased. Extremophile organisms contain enzymes that withstand hard conditions to survive (Tsigos *et al.*, 1998). In this context, *Thermoplasma acidophilum* (Darland *et al.*, 1970) (Ruepp *et al.*, 2000) is an archaea with a source of enzymes with robust features to survive in extreme conditions, as well as having the potential for biotechnological applications.

^{*} Corresponding author. E-mail: lsantos@ipicyt.edu.mx Tel. +52-444-8342057, Fax +52-444-8342010

Table 1. Closest structural homologues displayed by the protein data bank analyses

PDB Hit	Description	Oligomeric state	Ion dependence	TM-score	Reference
4JBG	Pyrobaculum aerophilum ADH	Tetramer	Zn ⁺²	0.964	(Vitale et al. 2013)
2EIH	Thermus thermophilus HB8 ADH	Tetramer	Zn ⁺²	0.959	To be published
4A0S	Streptomyces sp JS360 Octenoyl-CoA reductase/carboxylase	Tetramer	-	0.936	(Quade et al. 2012)
1RJW	Bacillus stearothermophilus strain LLD-R htADH	Tetramer	Zn ⁺²	0.931	(Ceccarelli et al. 2004)
3KRT	Streptomyces coelicolor A3(2) Crotonyl CoA reductase	Tetramer	Cl ⁻	0.930	To be published
3MEQ	Brucella melitensis ADH	Tetramer	Zn ⁺² , Cl ⁻ , Na ⁺	0.930	To be published
1R37	Sulfolobus solfataricus ADH	Tetramer	Zn^{+2}	0.929	(Esposito et al. 2003)
4WGZ	Saccharomyces cervisiae ADH	Tetramer	Zn ⁺²	0.928	(Raj et al. 2014)
3S1L	Cupriavidus necator JMP134 ADH	Tetramer	Zn ⁺²	0.927	(Kang et al. 2012)
2EER	Sulfolobus tokodaii strain7 ADH	Tetramer	Zn ⁺²	0.925	To be published

We have previously reported the analysis of the oxidative reaction of *T. acidophilum* NAD⁺ dependent Ta1316 ADH throughout the biocatalysis of ethanol to acetaldehyde (Marino-Marmolejo *et al.*, 2009). This work describes the enzymatic characterization towards the reductive reaction in the presence of ketones, aldehydes and keto-esters.

2 Materials and methods

2.1 Chemicals

All chemicals were analytical grade; aldehydes, ketones and keto-esters were purchased from Sigma-Aldrich.

2.2 Sequence analysis

Homologues of Ta1316 ADH (NCBI reference sequence: NP_394771.1) sequenced open reading frames were identified using the BLAST tool server at NCBI (www.blast.ncbi.nlm.nih.gov/). The 3D structure prediction was performed by using the I-TASSER server (Zhang, 2008; Roy *et al.*, 2010; Roy *et al.*, 2012) and the UCSF-Chimera program for interactive analyses of the structures (Meng *et al.*, 2006).

2.3 Expression and purification of the recombinant Ta1316 ADH

Ta1316 ADH cloned into the vector pET28a(+) was expressed in E. coli BL21 (DE3) pLysS from Novagen (Madison, WI) and purified as previously described (Marino-Marmolejo *et al.*, 2009).

2.4 Mass spectrometry

Recombinant Ta1316 ADH was identified by mass spectrometry analysis (LC-ESI-MS/MS) in the LINAN laboratory facility at the IPICYT. Data identification was carried out on the MS/MS spectra datasets using the MASCOT search algorithm (Matrix Science, London, UK). Searches were performed using the NCBI database.

2.5 Reduction assay

Consumption of NADH to NAD⁺ (oxidation) was measured at 340 nm (Kagi & Vallee, 1960) using a Variant Cary 50 spectrophotometer with a peltier temperature-controlled cuvette-holder and the Kinetics v3.0 software (Variant). The reaction mixture was completed in a total volume of 1 ml containing 50 mM phosphate buffer (pH 5.0), 1 mM NADH, 0.5 mM ZnSO₄, 10 μ g of pure recombinant protein, 2 mM pure aldehyde, or 1 mM aldehydes, ketones or

34

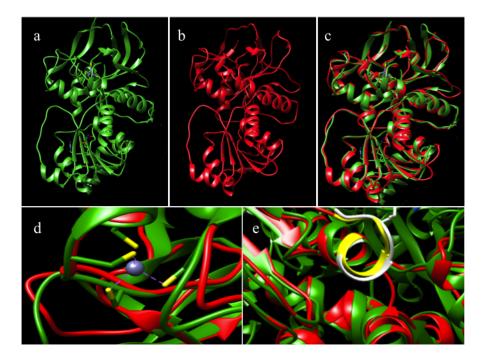


Fig. 1: In silico modeling of Ta1316 ADH against its closest homologue. a) Crystal structure of P. aerophilum str. IM2 ADH, PyAeADHIII (green), and b) 3D model of Ta1316 ADH (red). c) Superposition of PyAeADHIII (red) and Ta1316 ADH (red). d) The active ligand site is zoomed in where the Zn2+ ion is represented as a sphere. e) Sequences of the NAD-binding site in Ta1316 ADH and PyAeADHII are represented in yellow and white respectively.

keto-esters (Table 2). All pH, temperature, substrate specificity, kinetics and effects of compounds studies were conducted in triplicate. Maximum activity was determined with a temperature range of 25 to 90°C and pH range of 2 to 8 at 75°C. K_m and V_{max} values were estimated using the method of least squares.

3 Results

Recombinant Ta1316 ADH was identified by electrospray ionization mass spectrometry and MASCOT dataset. The average nominal mass reported of 36278 Da was equivalent to the calculated theoretical value of the sequence. 57% sequence coverage of matched peptides was confirmed,

Table 2. Eff	ect of	chemical	s on the	Ta1316	ADH	activity
--------------	--------	----------	----------	--------	-----	----------

Chemicals	Concentration (mM)	Relative activity (%) ^a
None	-	100.00
$ZnSO_4$	1	194.78
$MnCl_2$	1	130.96
$CoSO_4$	1	117.52
$MgCl_2$	1	95.55
FeCl ₃	1	73.71
$CuCl_2$	1	0.001
EDTA 2Na	10	68.63
2-Mercaptoethanol	1	46.41
N-Ethylmaleimide	1	52.89

^aThe activity was measured at 75°C as described in Materials and Methods section in the presence of 2 mM acetaldehyde

www.rmiq.org 35

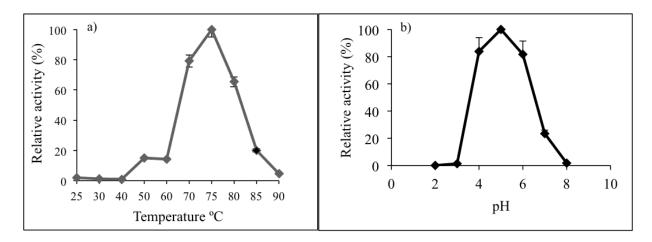


Fig. 2: a) Effect of temperature on Ta1316 ADH activity. b) Effect of pH on Ta1316 ADH activity.

and the NCBI Entrez gi 10640659 or 16082305 reports indicated an alcohol dehydrogenase from *Thermoplasma acidophilum* DSM 1728.

Structural analyses displayed by the PDB platform showed the crystal structure of *Pyrobaculum aerophilum* ADH (Vitale *et al.*, 2013) as the closest structural conformation against Ta1316 ADH (Figure 1, Table 1). Unexpectedly, *P. aerophilum* does not possess any apparent ADH activity for comparison against Ta1316 ADH. The second best TM-score found, *Thermus thermophilus* ADH (Pennacchio *et al.*, 2008), was similar and presents a comparable substrate preference.

The enzymatic oxidation reaction of NADH to NAD⁺ produced a reaction rate that increases up to 75°C as shown in Figure 2a, then activity decays gradually until inactivation at 90°C. The maximum temperature found is comparable to *Sulfolobus acidocaldarius* ADH (Pennacchio *et al.*, 2010).

Maximum activity displayed by Ta1316 ADH was at pH 5.0 in the presence of 2 mM acetaldehyde (Figure 2b). This pH value is closely related to the intracellular pH of *T. acidophilum* (pH 5.5) (Searcy, 1976) and those found for the reduction of NAD+ (Marino-Marmolejo *et al.*, 2009) and *S. acidocaldarius* (Pennacchio *et al.*, 2010).

Table 3. Substrate specificity of Ta1316 ADH in the reduction reaction

Substrate (1 mM)	Relative activity (%)	
Ketones		
3'-Chloroacetophenone	67.98	
3-Octanone	61.57	
4-Acetylpyridine	53.68	
2'-Nitroacetophenone	14.91	
Acetylacetone	0	
4-Fluoroacetophenone	0	
Aldehydes		
2-Chlorobenzaldehyde	60.06	
3-Nitrobenzaldehyde	59.10	
Benzaldehyde	58.07	
2-Nitrobenzaldehyde	42.61	
Keto-esters		
Methyl pyruvate*	100	

^{*}The activity towards methyl pyruvate (1 mM) was taken as 100%.

Table 4. Biochemical features of several ADH members in the reduction reaction

Source	Specific activity (U/mg)	optimum pH	optimum T(°C	Km (mM)	Kcat/Km (s ⁻ 1mM ⁻ 1)	Reference		
Ta1316 ADH ^{a,b}	ND	5.0	75	3.97	46	0.56	11.6	This work
Geobacillus	ND	8.0	60	4.71	404.7	568.7	85.94	(Liu et al. 2009)
thermodenitrificans NG80-2								
ADH1 ^a								
Desulfovibrio gigas ^a	31.8	7.5	40	1.8	ND	ND	ND	(Hensgens et al. 1993)
Thermococcus strain ES1 ^b	ND	7.0	ND	0.84	22.9	30.5	27	(Ying et al. 2009)
Mus musculus ^c A2	0.012	9.5	30	0.041	ND	17.83	ND	(Algar et al. 1983)
Rattus norvegicus ^c ADH3	ND	7.5	25	0.17	170	ND	1000	(Julia et al. 1987)

ND, Not determined; ^aArchaea; ^bBacteria; ^cEukaryote; ^d2mM acetaldehyde as substrate.

The effect of various divalent cations was also examined (Table 2). Apparently, no metal cofactor is required for the enzymatic activity however, CoSO₄, MnCl₂ and ZnSO₄ produced increasing activities of 117, 130 and 194% respectively against the blank reaction (no metal ion added). It has been reported that ZnCl2 is able to inhibit Oenococcus oeni or Yokenella sp. ADH activities (Elleuche et al., 2013) (Ying et al., 2014). Surprisingly, in the presence of MgCl₂, Ta1316 ADH conserved 95.5% activity. FeCl₃ retained virtually 75% activity. Moreover, evident obstruction of activity was observed in the presence of the Cu+2 ion as reported for Moraxella (Tsigos et al., 1998). The metal chelating agent EDTA was used at increased concentrations and arrest of activity was only observed at 10 mM producing less than 50% inhibition (63% relative activity). This behavior has been reported for the yeast ADH (Magonet et al., 1992). It is expected that Zn⁺² ions confer high stability to the reduction reaction as shown for other ADHs (Kawano et al., 2011; Rong Chen, 2011) (Pennacchio et al., 2010) (Kizaki et al., 2008). Lastly, inhibition of Ta1316 ADH activity by the thiol group 2-mercaptoethanol (Cheng & Lek, 1992) (Langeland et al., 1999) and N-ethylmaleimide reduced 50% the activity.

Ta1316 ADH substrate specificity was standardized reaching a maximum activity of 100% at 1 mM methyl pyruvate (Table 3). We observed that in the presence of side chains ketones, such as 3'-chloroacetophenone, 3-octanone, 4-acetylpyridine and 2'-nitroacetophenone the enzyme showed decreasing relative activities of 68 to 15%. Surprisingly, the enzyme did not show activity with ketones such as acetylacetone or 4'-fluoroacetophenone in contrast to *Lactobacillus kefir* ADH (Chen *et al.*, 2010).

Aldehyde substrates (benzaldehydes) were evaluated for conversion to their corresponding aromatic alcohols. We found minor differences in the relative activities: 2-chlorobenzaldehyde > 3-nitrobenzaldehyde > benzaldehyde > 2-nitrobenzaldehyde (60-42%) as shown in Table

3. Ta1316 ADH showed similar activities when comparison was performed against *Devosia riboflavina* ADH (Kizaki *et al.*, 2008), and higher activity than *Thermococcus guaymasensis* and *Moraxella* sp. ADHs (Ying & Ma, 2011) (Tsigos *et al.*, 1998).

Lastly, we found Ta1316 ADH was able to reduce keto-ester methyl pyruvate producing its highest activity (adjusted to 100%) and preferred substrate of reaction (table 3). This substrate has been used with low efficiency by *Candida maris* ADH (Kawano *et al.*, 2011), however it is capable of processing acetylacetone in contrast to Ta1316 ADH.

Kinetic parameters were measured in the presence of acetaldehyde. Ta1316 ADH displayed higher affinity for acetaldehyde (K_m 3.97 mM and V_{max} 0.56 mM/s) rather than ethanol (Marino-Marmolejo *et al.*, 2009), suggesting that Ta1316 ADH's main function is the reduction of aldehydes and ketones. A comparative analysis of several ADHs is shown in Table 4 to support biochemical and kinetic data.

Discussion

An enzymatic reduction of prochiral substrates was performed to characterize the reducing activity of Ta1316 ADH. In general, ADHs show some advantages for asymmetric catalysis against chemical catalysis; especially for challenging reactions where the purpose is the production of enantiopure substances.

Biotransformation is therefore key in industry; highly valuable products such as short chain alcohols (fine chemicals) are used as intermediates, e.g. (*R*)-2-butanol, which is 120 times more expensive than 2-butanone. In this context, Ta1316 ADH displayed broad substrate specificity both in oxidation and reduction reactions with special preference towards the reductive activity. Characterization of the enzymatic properties showed that Ta1316 ADH prefers NAD+ as cofactor and ZnSO₄ enhances 200% the relative activity. Moreover, Ta1316 ADH has

the ability to reduce methyl pyruvate as its principal substrate. The recombinant form of Ta1316 ADH can be produced in high quantities in *E. coli*, it is stable in a broad range of temperature and pH and has a wide substrate specificity in the reduction reaction. The properties reported here makes Ta1316 ADH an attractive catalyst in several industry processes, principally for the production of chiral compounds under extreme conditions.

Acknowledgments

Mabel Guzmán-Rodríguez was supported by the Graduate Scholarship No. 230931 from the National Council for Science and Technology (CONACYT), Mexico. We thank Mr. Graham Matthew Tippett for the English correction of the manuscript.

References

- Chen, Q., Hu, Y., Zhao, W., Zhu, C. y Zhu, B. (2010). Cloning, expression, and characterization of a novel (s)-specific alcohol dehydrogenase from lactobacillus kefir. *Applied Biochemistry Biotechnology 160*, 19-29.
- Cheng, L.Y. y Lek, L.H. (1992). Inhibition of alcohol dehydrogenases by thiol compounds. *FEBS Letters* 300, 251-253.
- Darland, G., Brock, T.D., Samsonoff, W. y Conti, S.F. (1970). A thermophilic, acidophilic mycoplasma isolated from a coal refuse pile. *Science* 170, 1416-1418.
- Eichler, J. (2001). Biotechnological uses of archaeal extremozymes. *Biotechnology Advances* 19, 261-278.
- Elleuche, S., Fodor, K., Klippel, B., von der Heyde, A., Wilmanns, M. y Antranikian, G. (2013). Structural and biochemical characterisation of a nad(+)-dependent alcohol dehydrogenase from *oenococcus oeni* as a new model molecule for industrial biotechnology applications. *Applied Microbiology Biotechnology 97*, 8963-8975.
- Kagi, J.H. y Vallee, B.L. (1960). The role of zinc in alcohol dehydrogenase. V. The effect of metal-binding agents on the structure of the yeast alcohol dehydrogenase molecule. *Journal* of Biological Chemistry 235, 3188-3192.

- Kawano, S., Yano, M., Hasegawa, J. y Yasohara, Y. (2011). Purification and characterization of an nadh-dependent alcohol dehydrogenase from candida maris for the synthesis of optically active 1-(pyridyl)ethanol derivatives. *Bioscience, Biotechnology, and Biochemistry* 75, 1055-1060.
- Kizaki, N., Yasohara, Y., Nagashima, N. y Hasegawa, J. (2008). Characterization of novel alcohol dehydrogenase of devosia riboflavina involved in stereoselective reduction of 3-pyrrolidinone derivatives. *Journal of Molecular Catalysis B:* Enzymatic 51, 73-80.
- Kroutil, W., Mang, H., Edegger, K. y Faber, K. (2004). Recent advances in the biocatalytic reduction of ketones and oxidation of secalcohols. Current Opinion in Chemical Biology 8, 120-126.
- Langeland, B.T., Morris, D.L. y McKinley-McKee, J.S. (1999). Metal binding properties of thiols; complexes with horse liver alcohol dehydrogenase. *Comparative Biochemistry and Physiology -Part B: Biochemistry and Molecular Biology 123*, 155-162.
- Magonet, E., Hayen, P., Delforge, D., Delaive, E. y Remacle, J. (1992). Importance of the structural zinc atom for the stability of yeast alcohol dehydrogenase. *Biochemical Journal 287 (Pt 2)*, 361-365.
- Marino-Marmolejo, E.N., De Leon-Rodriguez, A., de la Rosa, A.P. y Santos, L. (2009). Heterologous expression and characterization of an alcohol dehydrogenase from the archeon thermoplasma acidophilum. *Molecular Biotechnology* 42, 61-67.
- Meng, E.C., Pettersen, E.F., Couch, G.S., Huang, C.C. y Ferrin, T.E. (2006). Tools for integrated sequence-structure analysis with ucsf chimera. *BMC Bioinformatics* 7, 339.
- Nie, Y., Xu, Y., Mu, X.Q., Wang, H.Y., Yang, M. y Xiao, R. (2007). Purification, characterization, gene cloning, and expression of a novel alcohol dehydrogenase with anti-prelog stereospecificity from candida parapsilosis. *Applied Environmental Microbiology* 73, 3759-3764.

38

- Nie, Y., Xu, Y., Yang, M. y Mu, X.Q. (2007). A novel nadh-dependent carbonyl reductase with unusual stereoselectivity for (r)-specific reduction from an (s)-1-phenyl-1,2-ethanediol-producing micro-organism: Purification and characterization. *Letters in Applied Microbiology* 44, 555-562.
- Pennacchio, A., Giordano, A., Pucci, B., Rossi, M. y Raia, C.A. (2010). Biochemical characterization of a recombinant short-chain nad(h)-dependent dehydrogenase/reductase from sulfolobus acidocaldarius. *Extremophiles* 14, 193-204.
- Pennacchio, A., Pucci, B., Secundo, F., La Cara, F., Rossi, M. y Raia, C.A. (2008). Purification and characterization of a novel recombinant highly enantioselective short-chain nad(h)-dependent alcohol dehydrogenase from thermus thermophilus. *Applied Environmental Microbiology* 74, 3949-3958.
- Reid, M.F. y Fewson, C.A. (1994). Molecular characterization of microbial alcohol dehydrogenases. *Critical Reviews in Microbiology* 20, 13-56.
- Reyes-Valadez, J.N., Quintana-Hernández P.A., Coronado-Velasco, C, Castro-Montoya, A.J. (2016). Simulation of the bioethanol production process from a mixture of glucose/xylose including temperature, ph and sugar concentration effects. Revista Mexicana de Ingeniería Química 15, 1-9.
- Rong Chen, X.Y., Chunna Yu, Qiuyan Wang, Xiaori Zhan and Tian Xie. (2011). Characterization of the recombinant group 3 alcohol dehydrogenase from thermotoga lettingae tmo. *African Journal of Microbiology Research* 5(23), 4004-4012.
- Roy, A., Kucukural, A. y Zhang, Y. (2010). Itasser: A unified platform for automated protein structure and function prediction. *Nature Protocols* 5, 725-738.
- Roy, A., Yang, J. y Zhang, Y. (2012). Cofactor: An accurate comparative algorithm for structure-based protein function annotation. *Nucleic Acids Research* 40, W471-477.
- Ruepp, A., Graml, W., Santos-Martinez, M.L., Koretke, K.K., Volker, C., Mewes, H.W., Frishman, D., Stocker, S., Lupas, A.N.

- y Baumeister, W. (2000). The genome sequence of the thermoacidophilic scavenger thermoplasma acidophilum. *Nature 407*, 508-513
- Searcy, D.G. (1976). Thermoplasma acidophilum: Intracellular ph and potassium concentration. *Biochimica Biophysica Acta* 451, 278-286.
- Tsigos, I., Velonia, K., Smonou, I. y Bouriotis, V. (1998). Purification and characterization of an alcohol dehydrogenase from the antarctic psychrophile moraxella sp. Tae123. *European Journal of Biochemistry* 254, 356-362.
- Vitale, A., Thorne, N., Lovell, S., Battaile, K.P., Hu, X., Shen, M., D'Auria, S. y Auld, D.S. (2013). Physicochemical characterization of a thermostable alcohol dehydrogenase from pyrobaculum aerophilum. *PLoS One* 8, e63828.
- Yakushi, T. y Matsushita, K. (2010). Alcohol dehydrogenase of acetic acid bacteria: Structure, mode of action, and applications in biotechnology. *Applied Microbiology Biotechnology* 86, 1257-1265.
- Yang, C., Ying, X., Yu, M., Zhang, Y., Xiong, B., Song, Q. y Wang, Z. (2012). Towards the discovery of alcohol dehydrogenases: Nad(p)h fluorescence-based screening and characterization of the newly isolated rhodococcus erythropolis wz010 in the preparation of chiral aryl secondary alcohols. *Journal of Industrial Microbiology & Biotechnology* 39, 1431-1443.
- Ying, X. y Ma, K. (2011). Characterization of a zinc-containing alcohol dehydrogenase with stereoselectivity from the hyperthermophilic archaeon thermococcus guaymasensis. *Journal of Bacteriology* 193, 3009-3019.
- Ying, X., Wang, Y., Xiong, B., Wu, T., Xie, L., Yu, M. y Wang, Z. (2014). Characterization of an allylic/benzyl alcohol dehydrogenase from yokenella sp. Strain wzy002, an organism potentially useful for the synthesis of alpha,beta-unsaturated alcohols from allylic aldehydes and ketones. *Applied Environmental Microbiology* 80, 2399-2409.
- Zhang, Y. (2008). I-tasser server for protein 3d structure prediction. *BMC Bioinformatics* 9, 40.