



Potential of microorganisms to decrease the "beany" off-flavor

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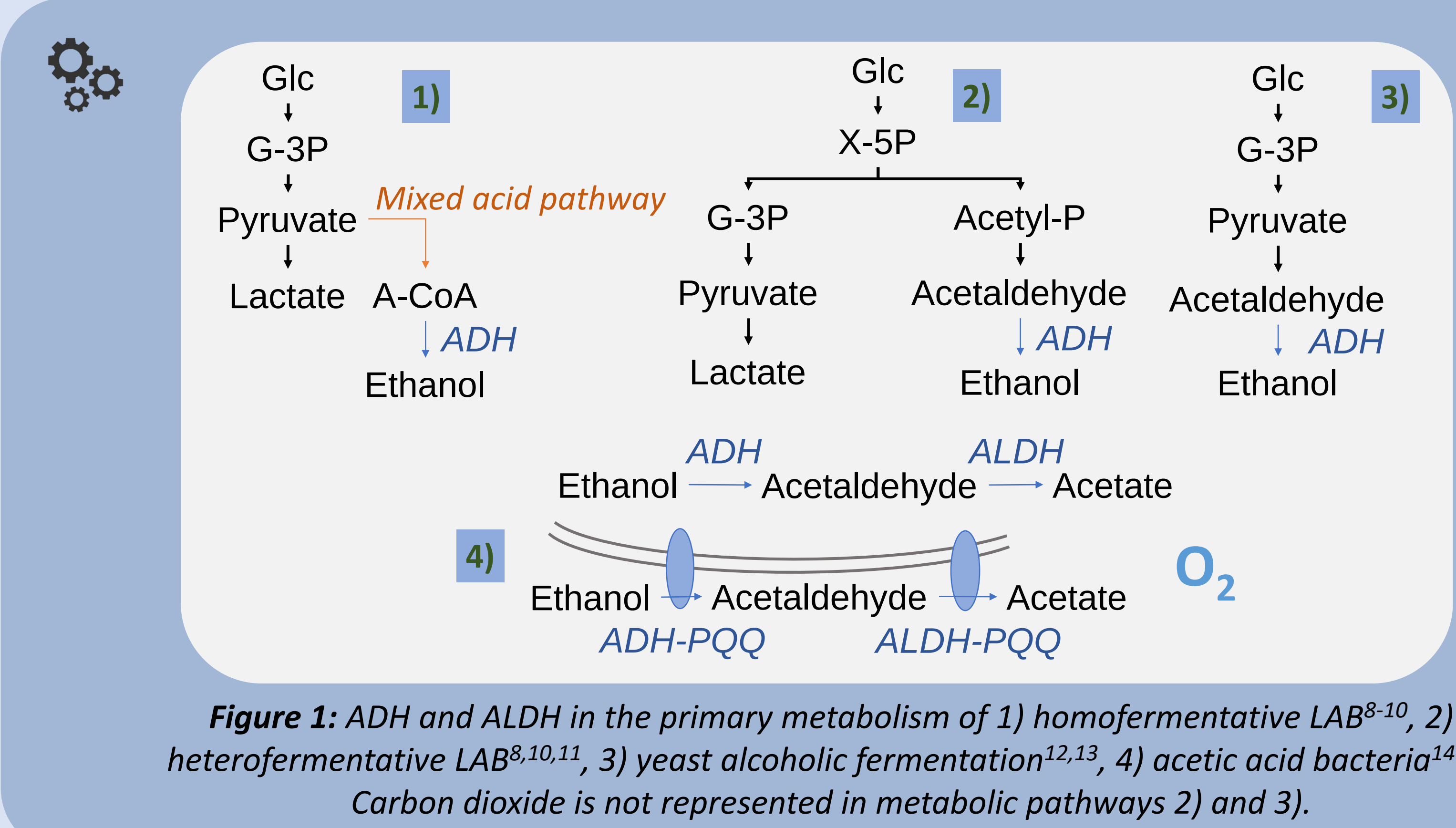
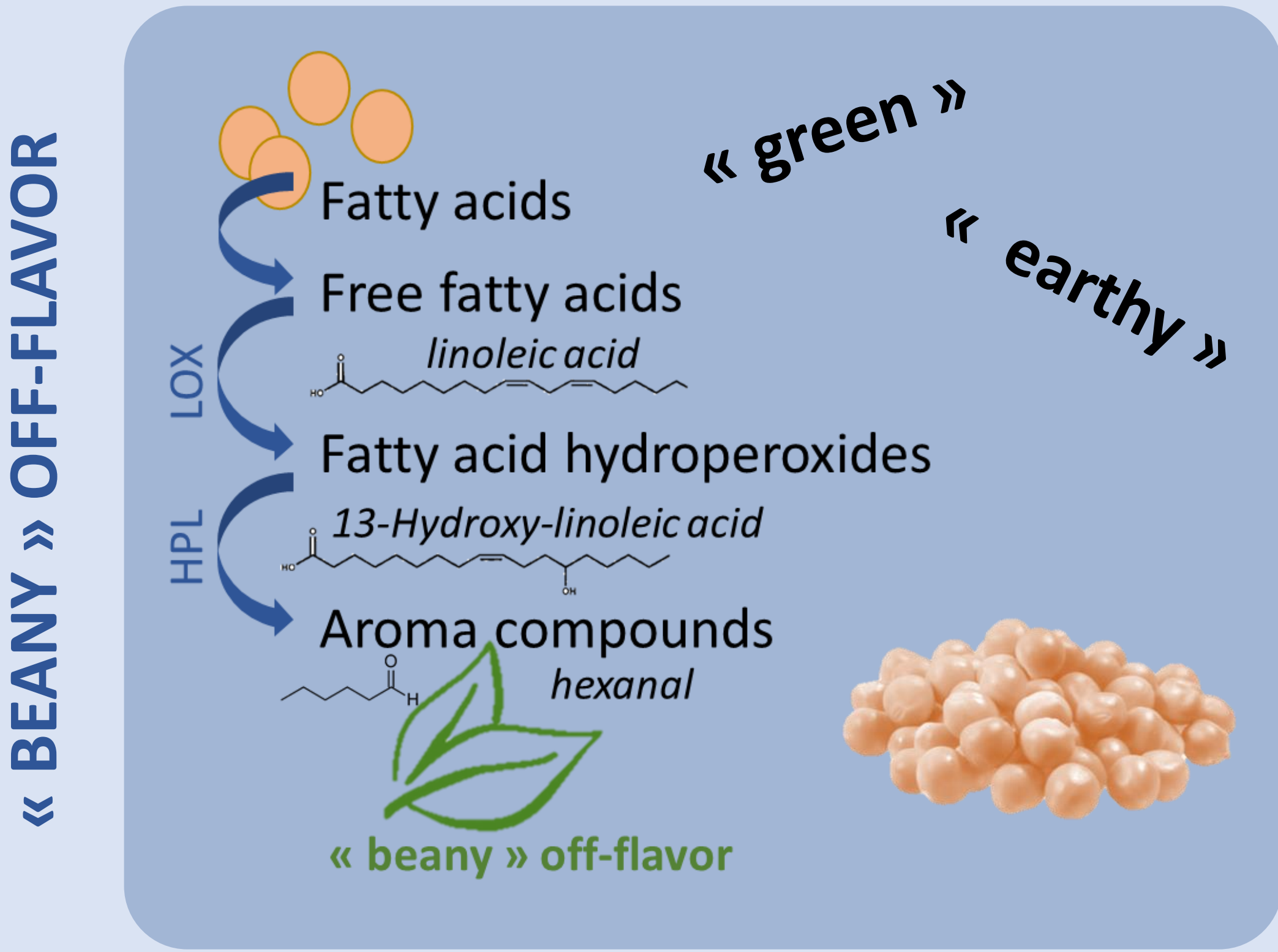
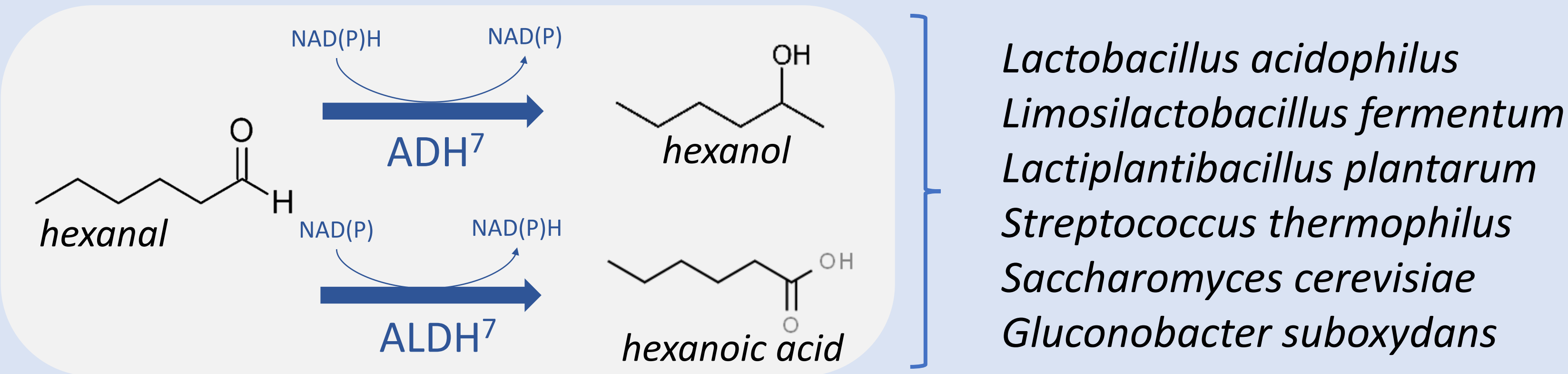
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POTENTIAL OF MICROORGANISMS TO DECREASE THE « BEANY » OFF-FLAVOR

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Plant-based protein consumption¹ → disadvantages : “beany” off-flavors².
Decrease the off-flavor ? Breeding, harvesting and storage, chemical and physical treatments.

Fermentation - Investigate the potential of mo to decrease the “beany”³⁻⁶



Involved ADH and ALDH in the strains catabolism (Fig. 1.)

Inductible or constitutive enzymes

Homo and HeteF inducible enzymes
HeteO, Y and AAB constitutive enzymes

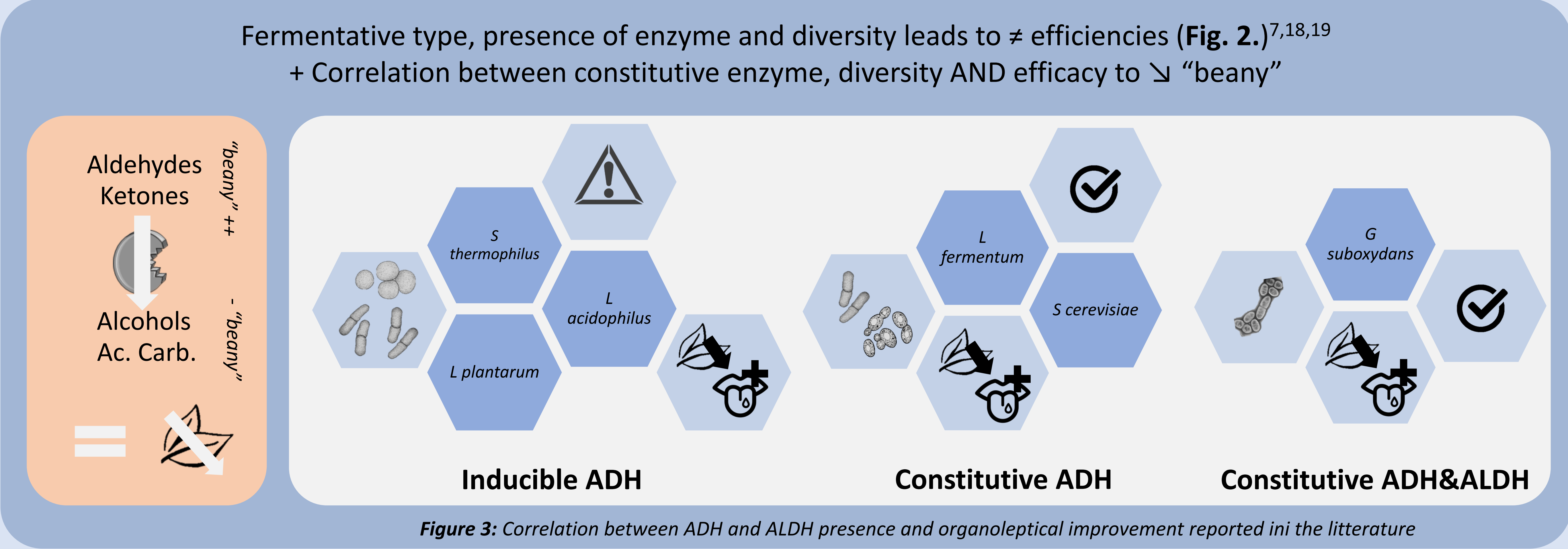
ADH and ALDH diversity, presence in the genome (Fig. 2.) KEGG¹⁷

Table 2: Bioinformatics data on ADH and ALDH

Species	ADH	ALDH
<i>L acidophilus</i>	3	6
<i>L fermentum</i>	4	8
<i>L plantarum</i>	4	11
<i>S thermophilus</i>	2	6
<i>S cerevisiae</i>	5	10
<i>G suboxydans</i>	4	13

Differences in enzyme diversity

≥ 4 ADH in He, Y and AAB
≥ 8 ALDH in He, Y and AAB



Constitutive enzyme = higher efficacy
LAB: ADH predominant, HeteO ++
Y: ADH++ ; AAB: ADH&ALDH++ polyvalent
ADH&ALDH activities correlated with sensory improvement

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FISCHER E., CAYOT N., CACHON R., (2022). Potential of microorganisms to decrease the « beany » off-flavor: a Review ; JAFc XX