

Analysis of soil microbial communities

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Methods of analyses of soil microbial communities

- **Cultivation** – determination of Colony Forming Units
 - + covers viable microorganisms
 - only 1-2% of soil microbes cultivable
- **Activity** – **determination of microbial activities**
 - enzyme activities, respiration, production / consumption of chemicals...
 - + covers viable microorganisms
 - dependent on conditions

Methods of analyses of soil microbial communities

- **Genetic analyses** – extraction of DNA / RNA, sequencing, comparison with databases
 - + detailed information (taxonomy, abundance of taxons, metabolic potential, transcriptomics – expressed genes, stress genes...)
 - costly and not as spread equipment
 - sometimes too detailed data (limited database data, laborious evaluation)

Methods of analyses of soil microbial communities

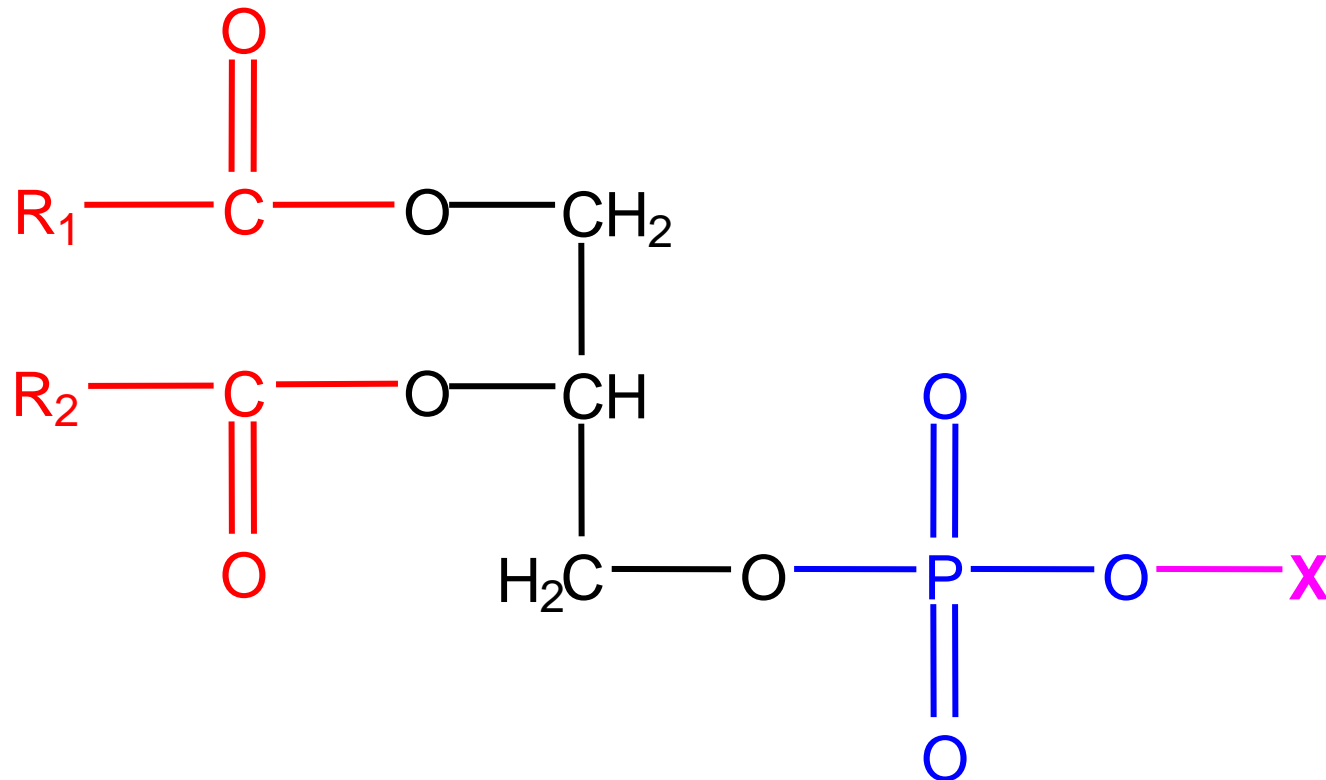
- **Chemical analyses** – determination of biomarker molecules / profiles
 - + in general simpler
 - + widespread and cheaper equipment
 - not as detailed information
 - possible interferences – need of careful interpretation



Biomarker molecules

- **Respiration isoprenoids** – chinons, length of side chain
- **Polysacharides** – surface, sheat
- **Polyamines**
- **Mycolic acids**
- **Sterols** – eucaryotic membrane (ergosterole in fungi)
- **Membrane lipids** – especially phospholipid fatty acid profiles
- ...

Phospholipids



R₁, R₂ – fatty acid acyls

X – hydrophilic groups

Phospholipids

- *In vivo* in membranes only
 - **cytoplasmatic**
 - outer membrane of G- bacteria
 - inner membranes of eukaryotes
- **Never storage compounds** → ~proportional content to biomass
- **Fast decomposition after cell death** → estimation of living biomass

Phospholipid fatty acids (PLFA)

- Composition of membrane PLFA depends on
 - species (taxonomy)
 - temperature
 - physiological state (stress detection)
 - nutrition

Utilization of PLFA analyses in soil ecology

- Quantification of **living** microbial biomass (total PLFA)
 - fast decomposition after cell death – **non-cultivation technique**
- Quantification of microbial groups
 - fungi / bacteria ratio
 - G+/G- ratio
 - abundance of other groups (actinobacteria, methanogenes, anaerobes...)
- **Monitoring of stress and soil disturbance**



Basic PLFA extraction

- **Extraction of total lipids**
 - **single-phase** mixture MeOH+chloroform+phosphate buffer
- **Separation of lipid fractions**
 - solid-phase extraction – silica columns
- **Derivatization**
 - Usually alkaline methanolysis
- **GC-MS**
 - usually polar column

Community biomarker fatty acids

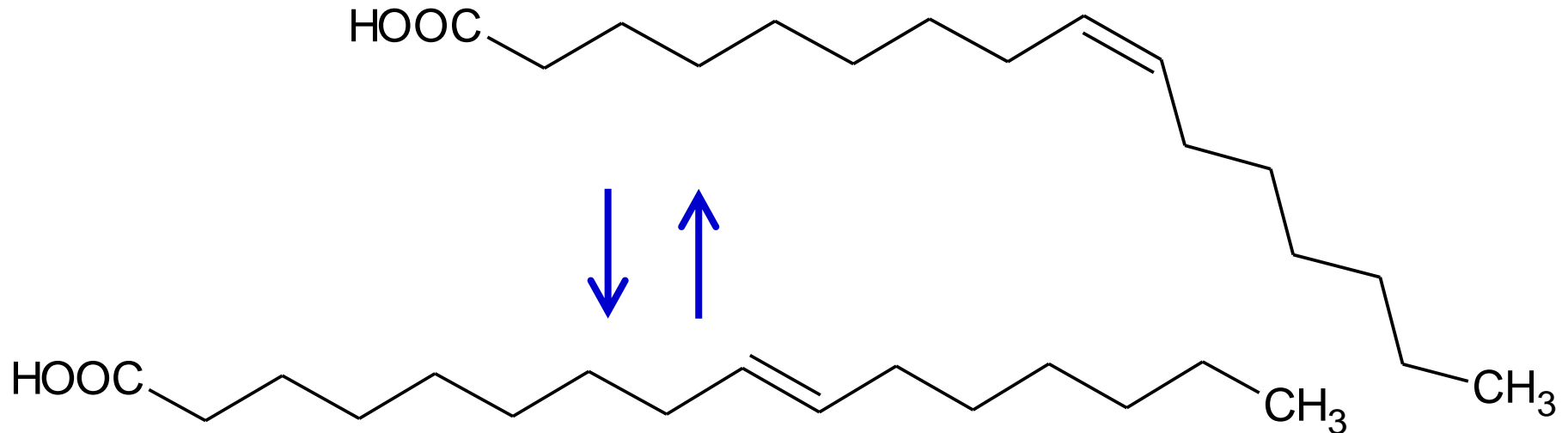
| Group | subgroup | Biomarker fatty acids |
|----------|----------------|---|
| Bacteria | G+ | i14:0, i15:0, a15:0, i17:0, a17:0 |
| | G- | cy17:0, cy19:0, 18:1w7 |
| | Actinobacteria | 10Me-16:0, 10Me-17:0, 10Me-18:0 |
| | Other | 16:1 ω 7t, 16:1 ω 7, 16:1 ω 9, |
| Fungi | | 18:2 ω 6,9 |
| Protozoa | | 20:4 ω 6 |

Physiological indicators

- Biochemical + empiric knowledge
- Only changes affecting membranes
- A series of published variants

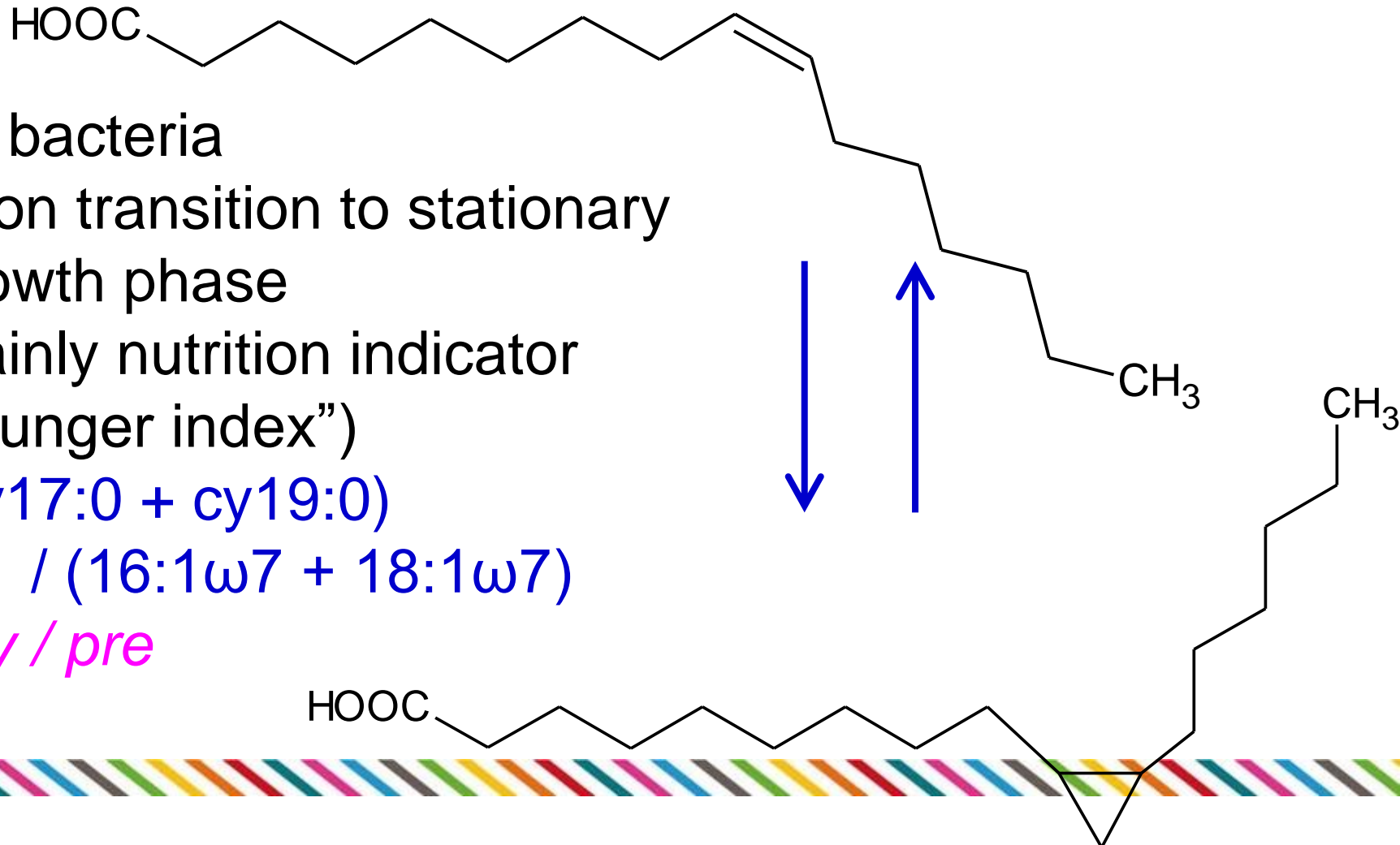


cis / *trans* isomeration



- Bacteria
- Changes directly in membrane
- *trans* / *cis* index
- general stress indicator
- $(18:1\omega7+16:1\omega7) / (16:1\omega7t+18:1\omega7t)$
- $>0.1 \rightarrow$ soil disturbance and stress

Cyclization of monounsaturated FA



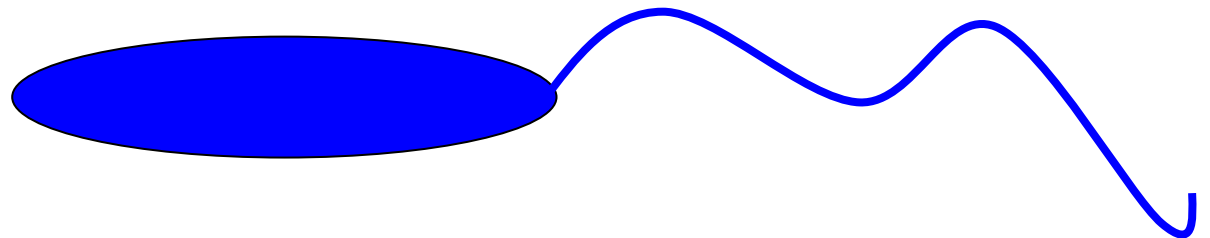
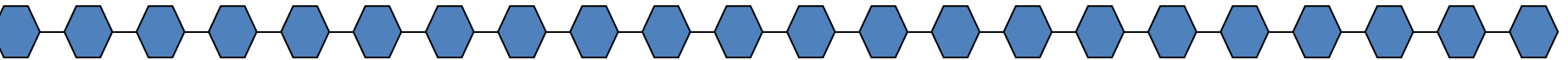
- G- bacteria
- upon transition to stationary growth phase
- mainly nutrition indicator (“hunger index”)
- (cy17:0 + cy19:0) / (16:1 ω 7 + 18:1 ω 7)
- *cy / pre*

Determination of microbial activities

- Completes biomass data with indication of activities
 - comparison of living biomass vers. activities gives useful information about overall state of community
- **Activity of extracellular enzymes** – decomposition of polymers
- **Respiration** – analyses of O₂ consumption or CO₂ production

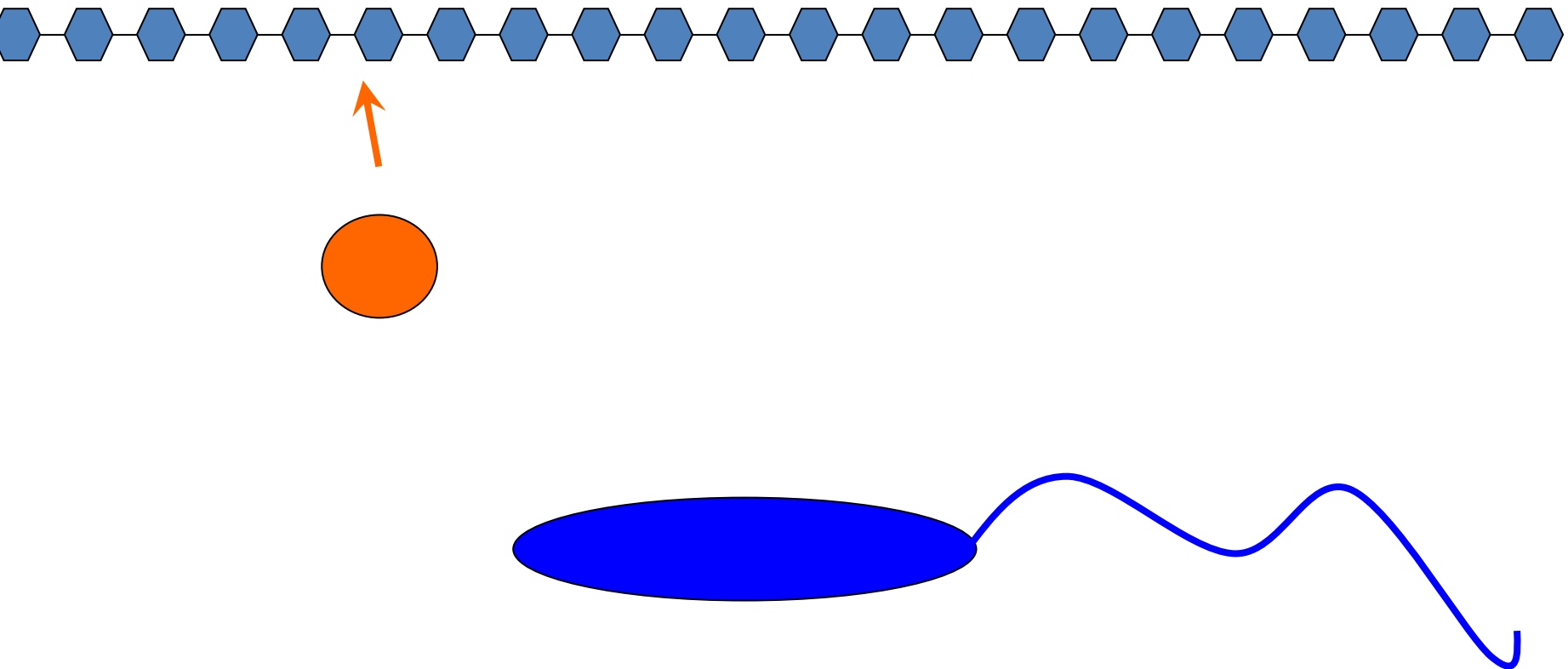
Determination of microbial activities

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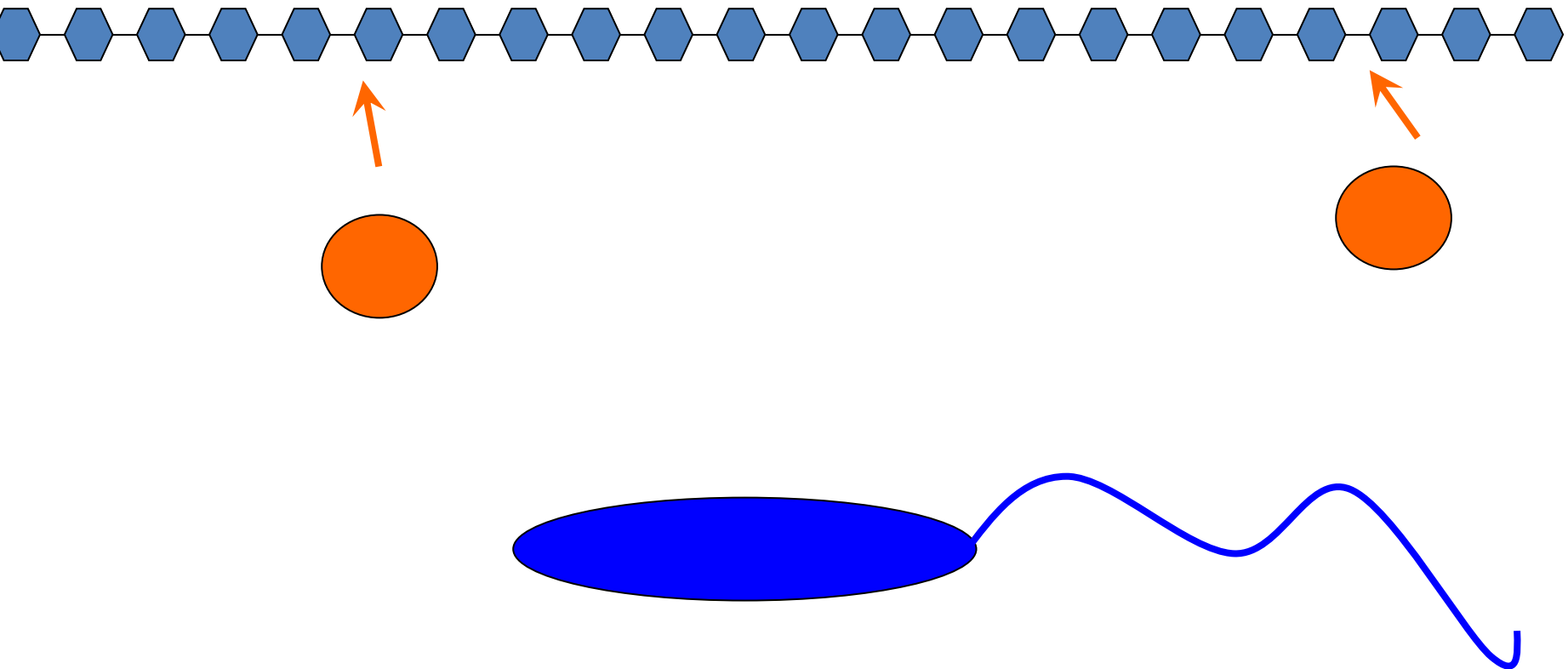
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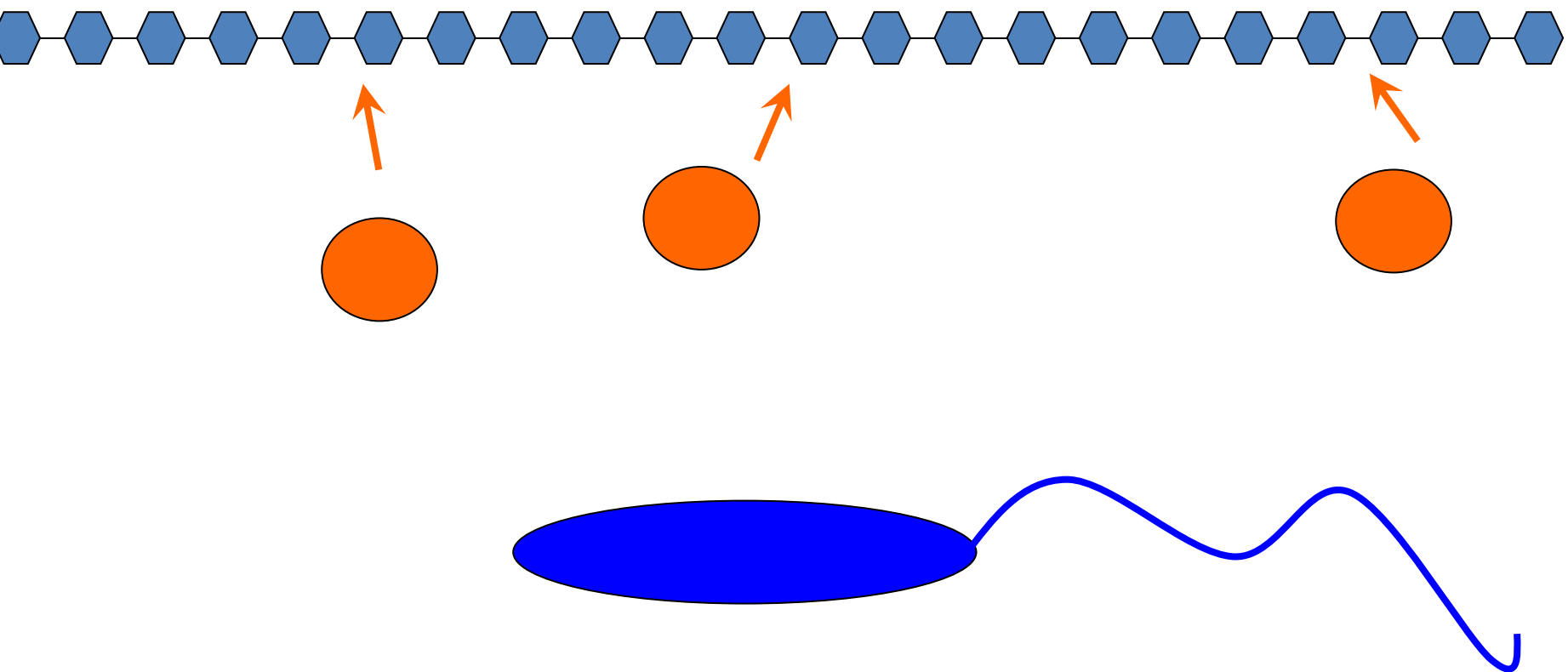
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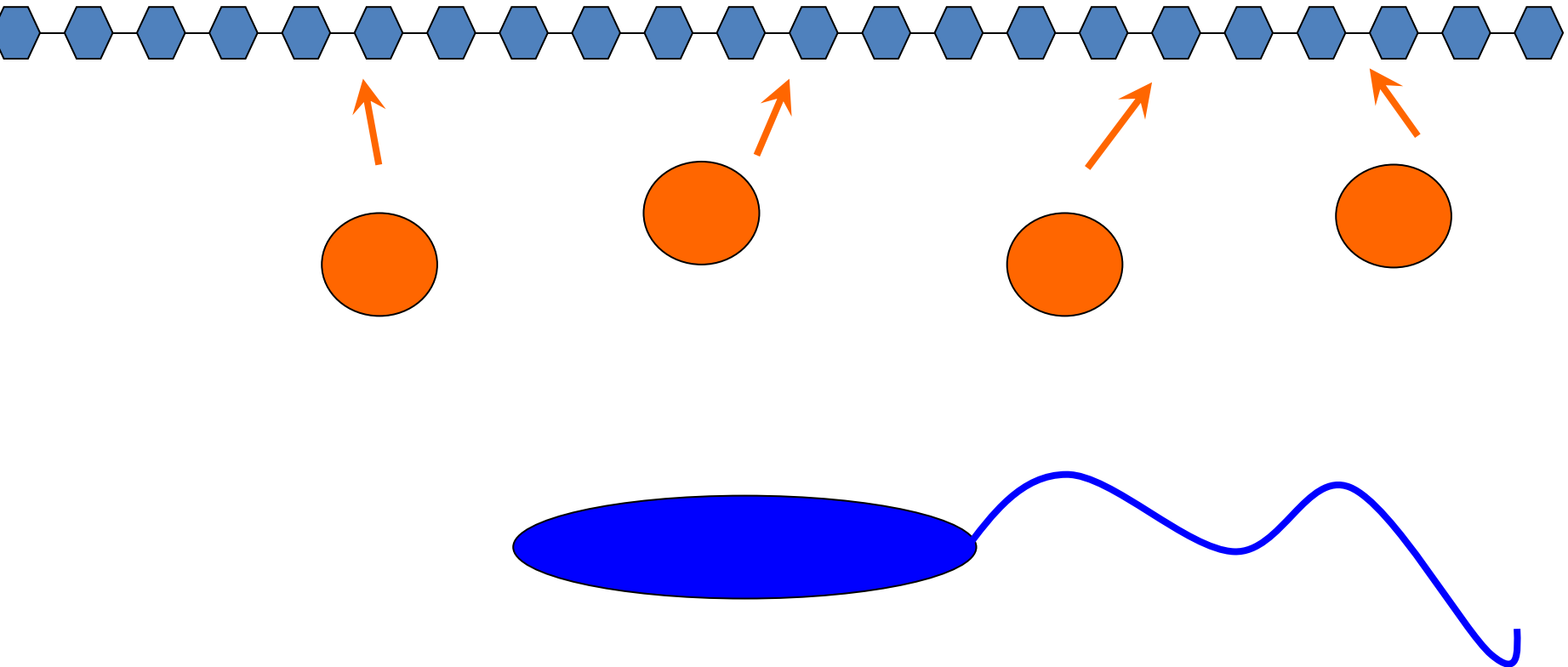
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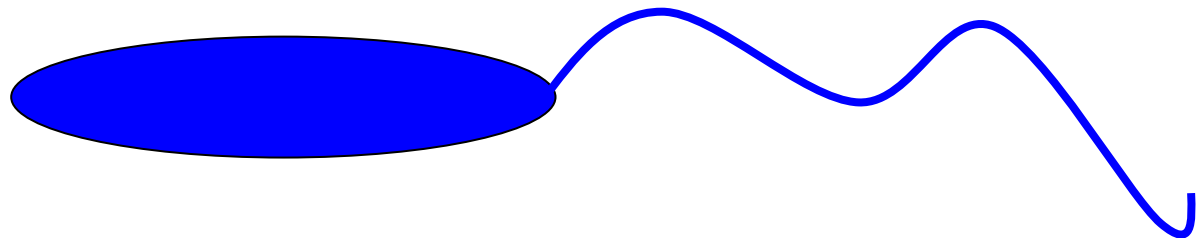
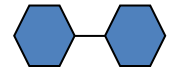
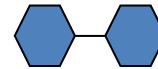
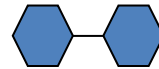
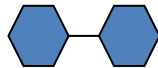
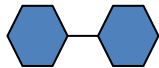
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Determination of microbial activities

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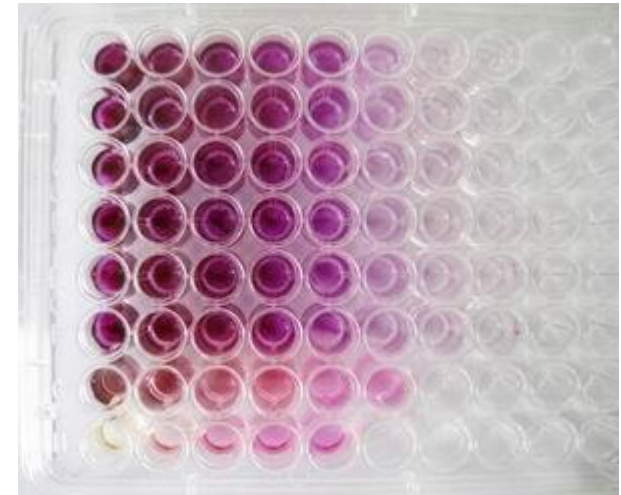
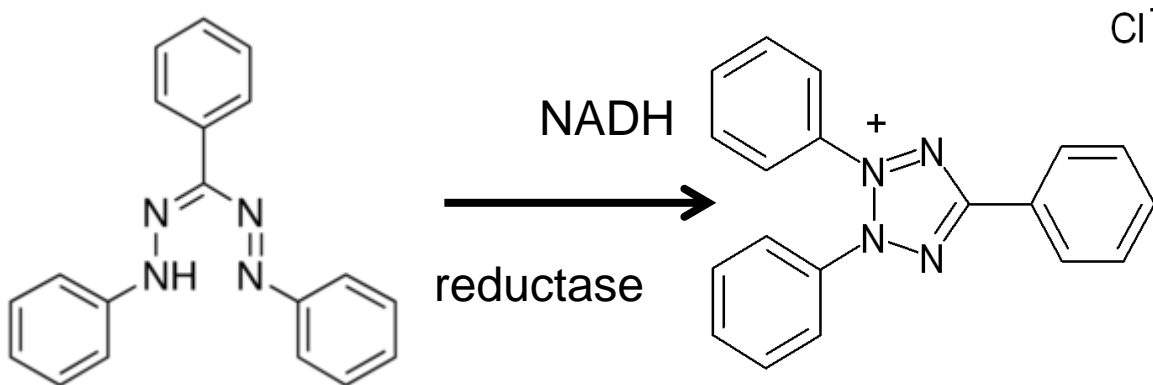
Activity of enzymes

- **Artificial enzyme substrates** – change to easily determined compounds
- **Many assays** – spectrophotometric or fluorometric
- High / low specificity



Tetrazolium assay

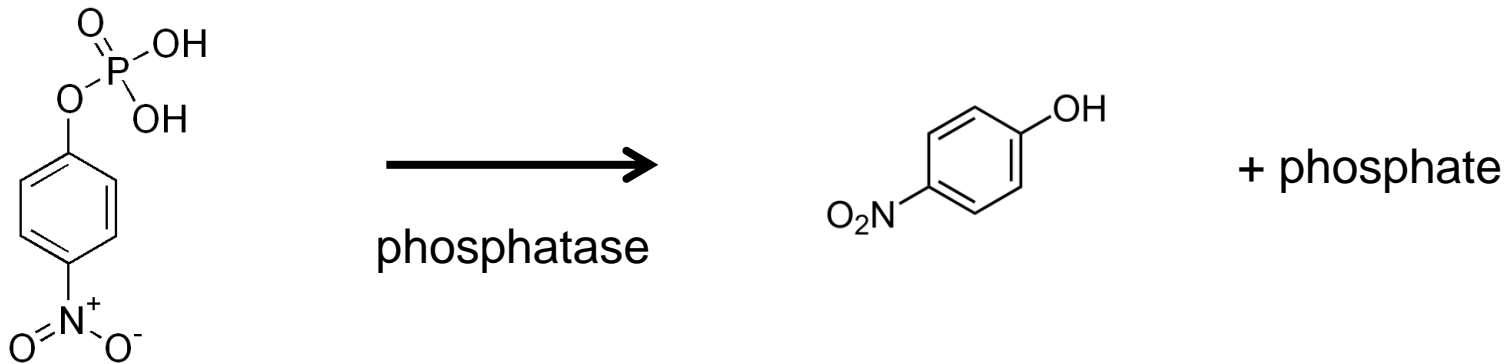
- Based on reduction of triphenyltetrazolium to triphenylphormazan
 - pink – spectrophotometric determination at 546 nm)
- Substrate of many oxidoreductases



- determines overall activity of soil or sludge

pNP assay

- Hydrolysis of pNP-X to p-nitrophenyl phosphate and X
- Determination of **yellow** pNP (pH >7, 400 nm)
- **Many variants** – phosphatases, sulphatases, proteases, glucosidases, chitinases...



Respiration

- Determination of the rate of
 - O₂ consumption
 - CO₂ production
- Simple titration
 - $\text{CO}_2 + \text{OH}^- \rightarrow \text{HCO}_3^-$
- Advance respirometers

- Variants:
 - optimization of humidity
 - addition of substrate (glucose, polysacharides...)