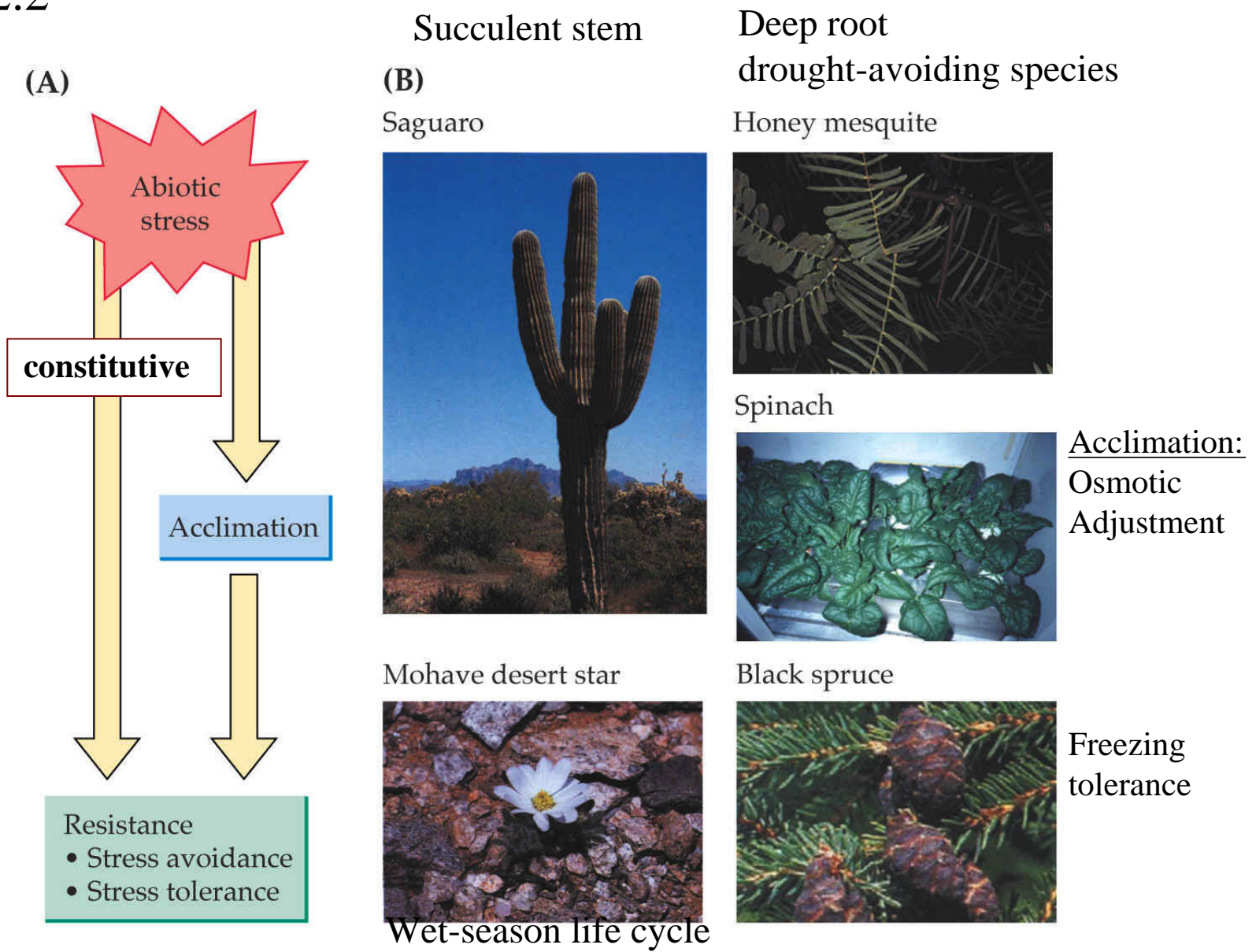


Fig. 22.2



Read 1158-1177

Many environmental conditions can lead to **water deficit**

-drought

-saline habitat

-low temperature (cold)

because water leaves cell and forms ice crystals in intercellular spaces

-transpirational water loss

$$U_w = U_s + U_p$$

U_w

water potential

U_s

solute potential, higher salt conc., lower the U_s

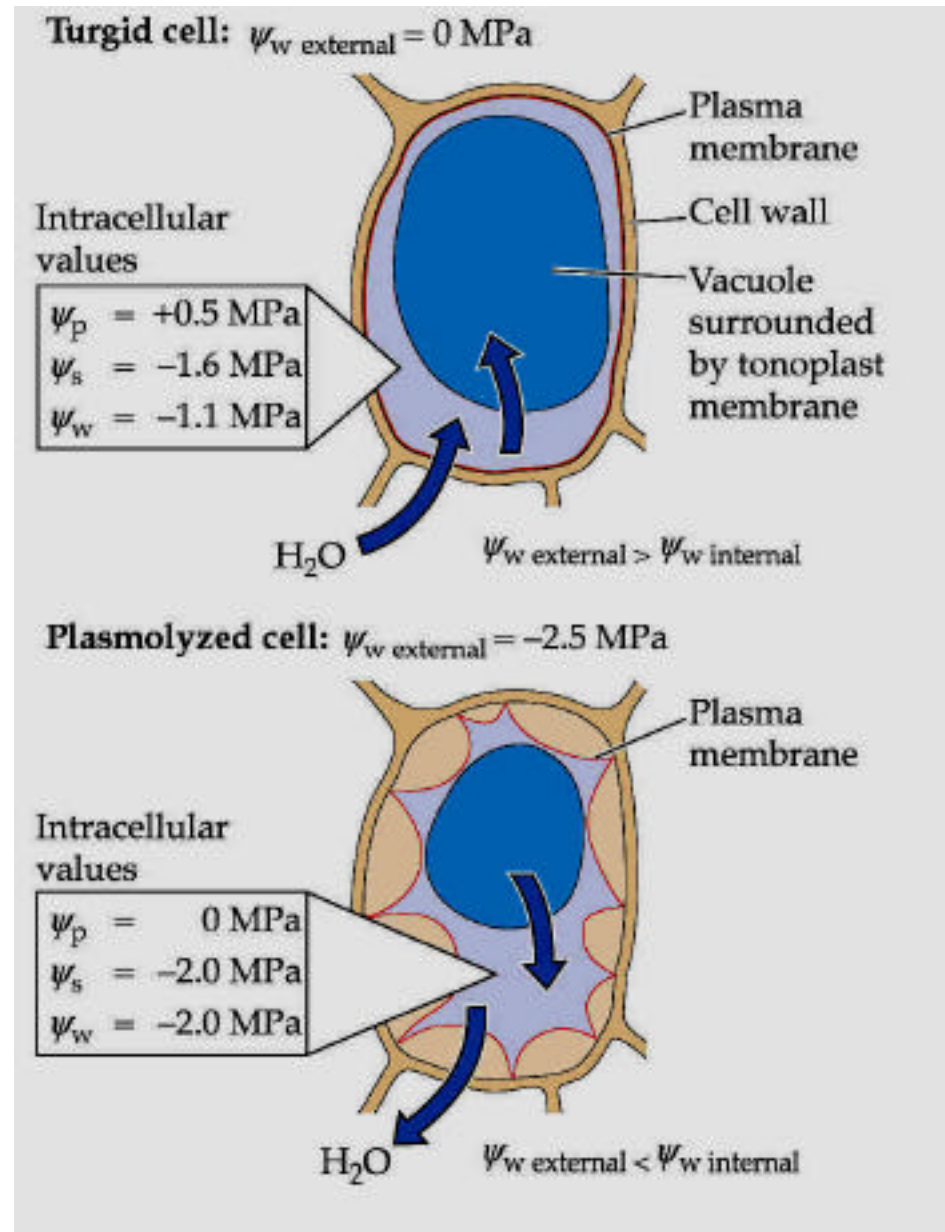
U_p

pressure potential,

physical force exerted on water to its environment

Water moves from high U_w to low U_w

Fig. 22.4



lethal

Osmotic adjustment

Increase in number of solute particles in a cell

Critical role in helping plants to acclimate to drought or saline conditions

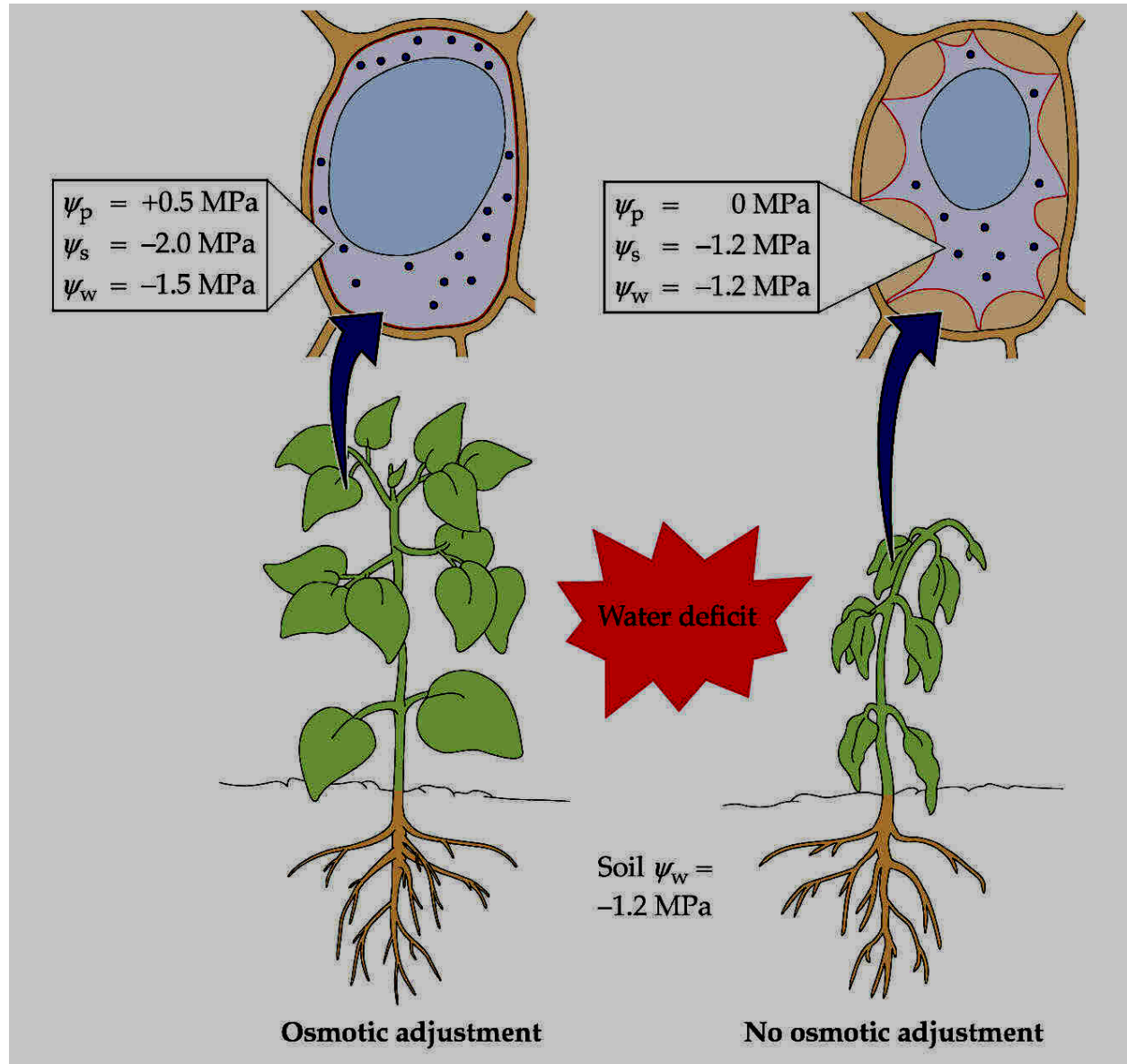
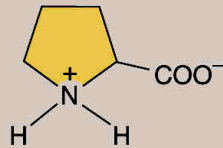


Fig. 22.6 Compatible osmolytes

Organic compound, that is highly soluble and non-toxic

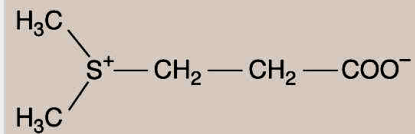
Compatible osmolytes

Amino acid:



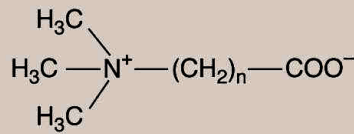
Proline

Tertiary sulfonium compound:

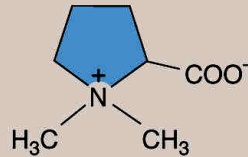


Dimethylsulfoniopropionate

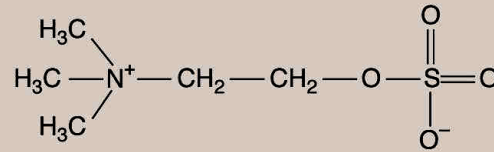
Quaternary ammonium compounds:



$n = 1$, Glycine betaine
 $n = 2$, β -Alanine betaine

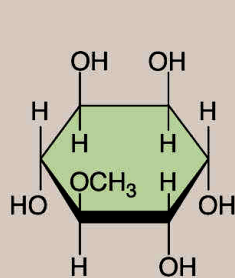


Proline betaine

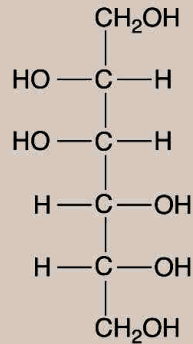


Choline-O-sulfate

Polyhydric alcohols:



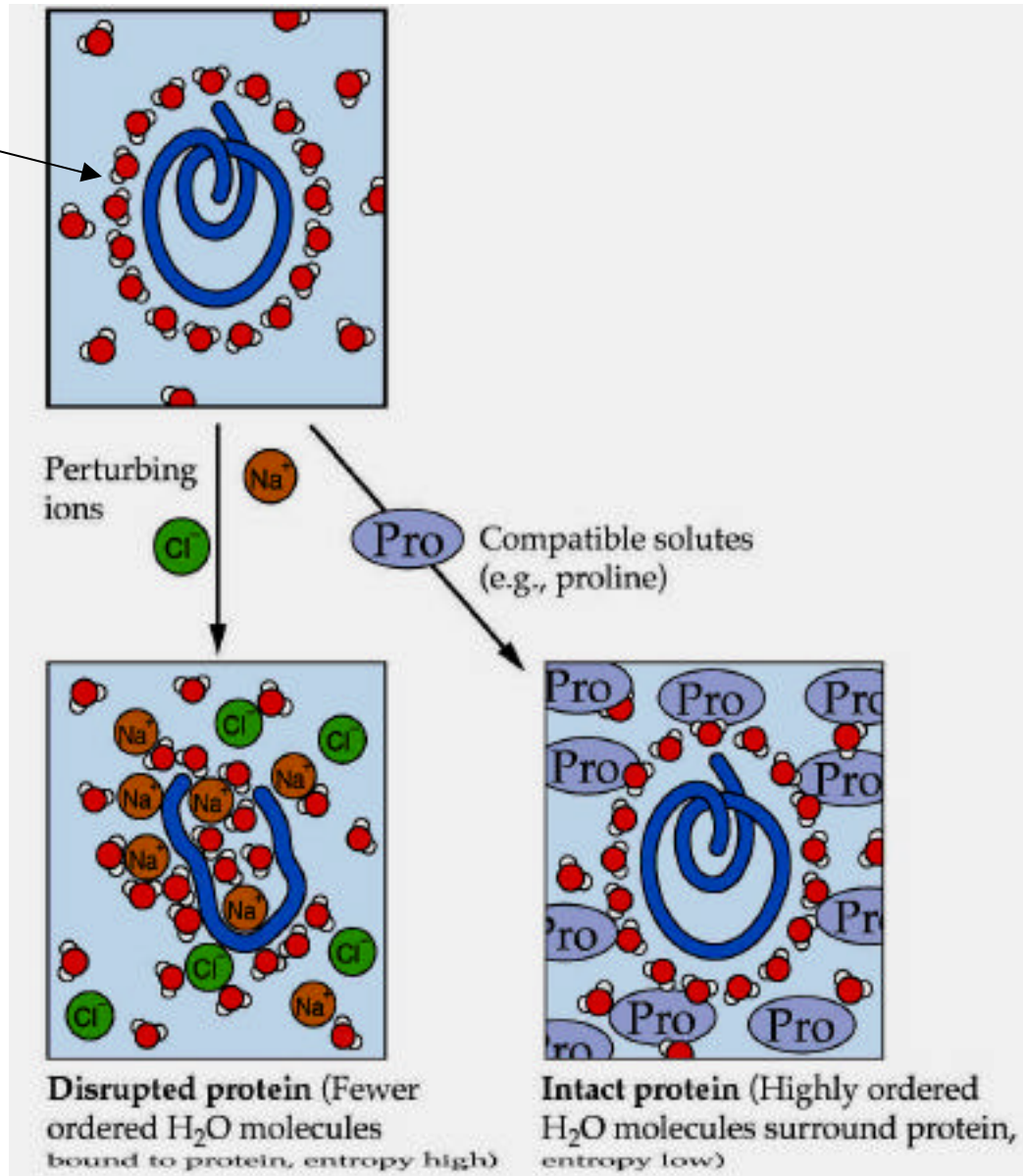
Pinitol



Mannitol

Fig. 22.7

hydration shell



Distribution vary in order to support osmotic equilibrium
among membrane bound organelles

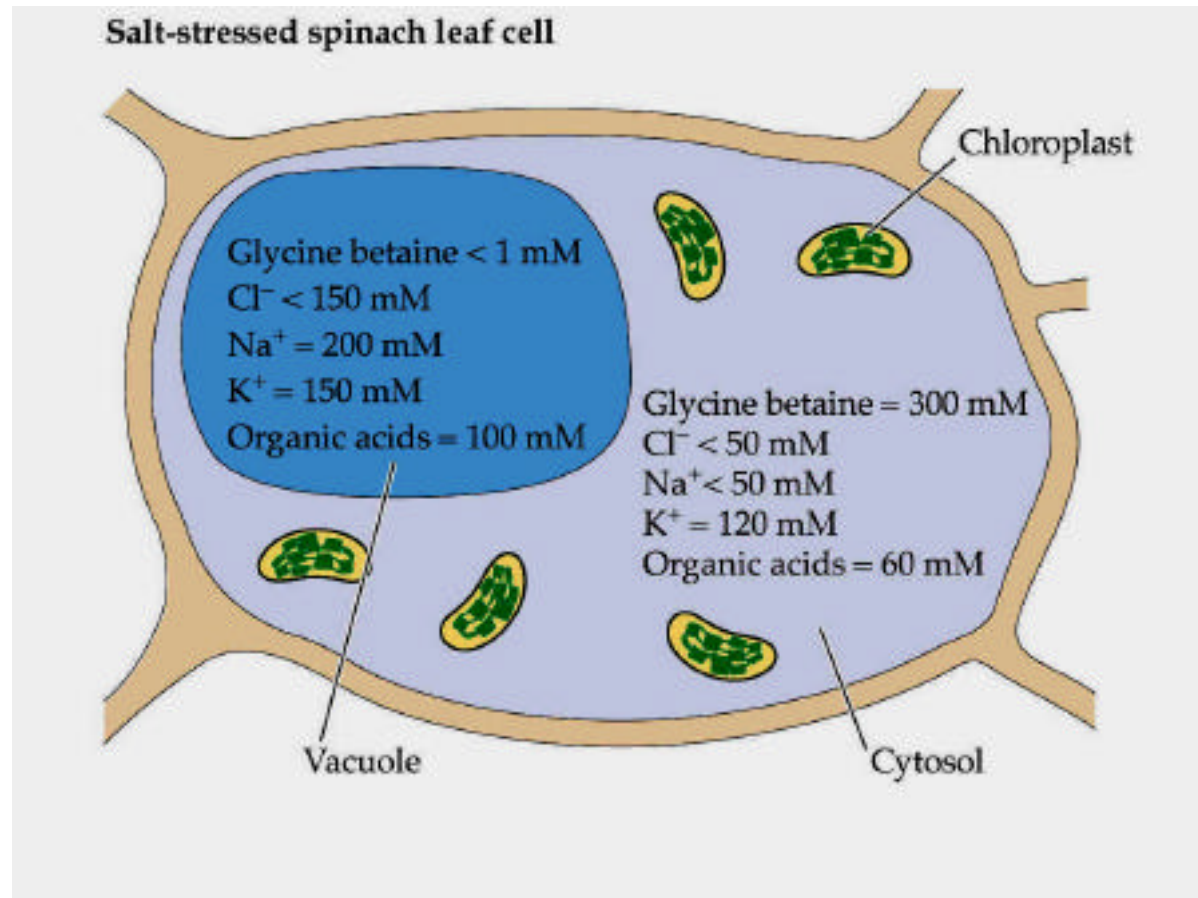


Fig. 22.8 spinach, high cytosolic glycine betaine allows the cytosol to achieve osmotic balance with vacuole which contains toxic ion and solutes

Genetic engineering of drought-tolerant plants

-Introducing biosynthetic enzyme for osmolytes

i.e. tobacco and Arabidopsis expressing mannitol-1-phosphate are more salt-tolerant

-Over-expression of Na⁺/H⁺ antiporter

Arabidopsis with such transgene can grow at 200mM NaCl (50% of sea salt conc.)

- Change of aquaporin gene expression

High Na⁺

Receptor

Ca⁺⁺ increase

SOS3 (Ca⁺⁺ binding)

SOS2 (Ser/Thr Kinase)

transporter (SOS1&?)
gene expression

nucleus

SOS1

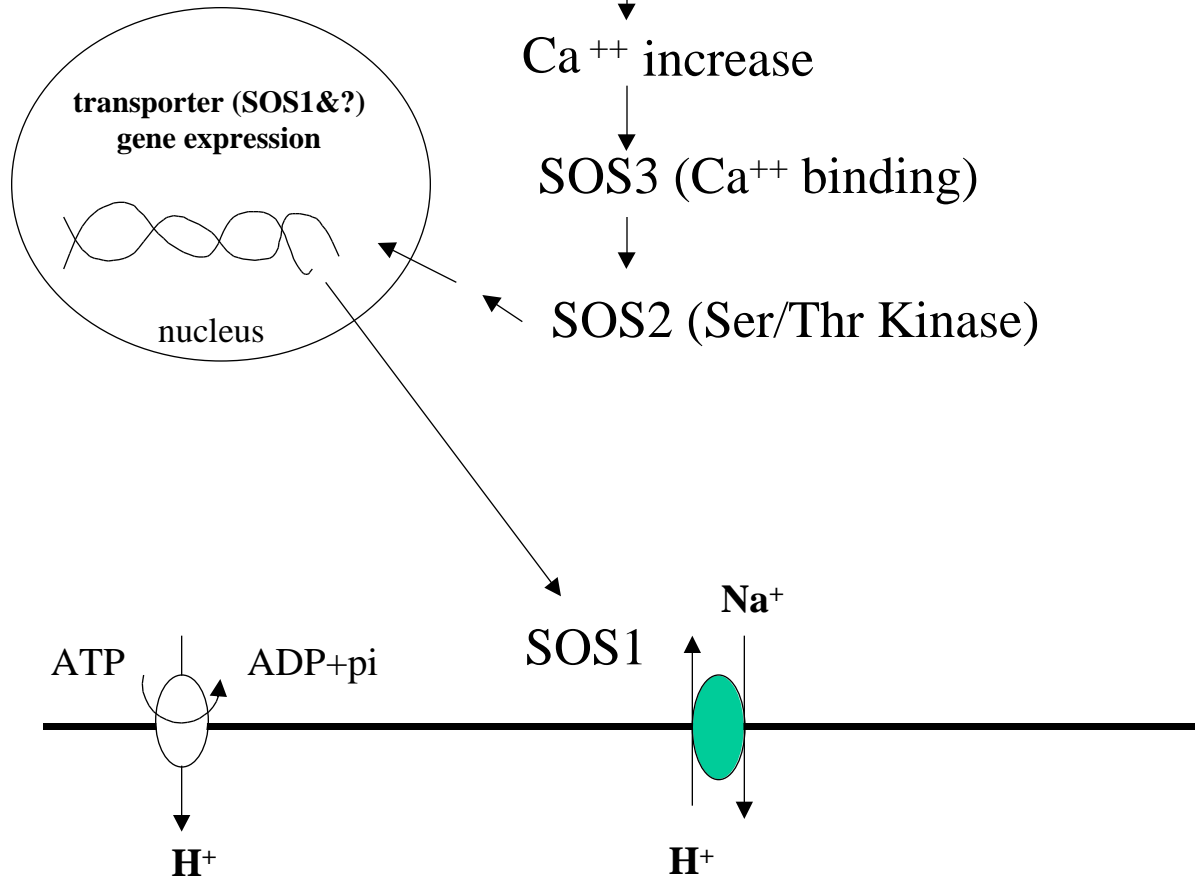
ATP

ADP+pi

Na⁺

H⁺

H⁺



Aquaporin: water channels

Members of MIP (major intrinsic protein) family located in plasma membrane

Fig. 3.52

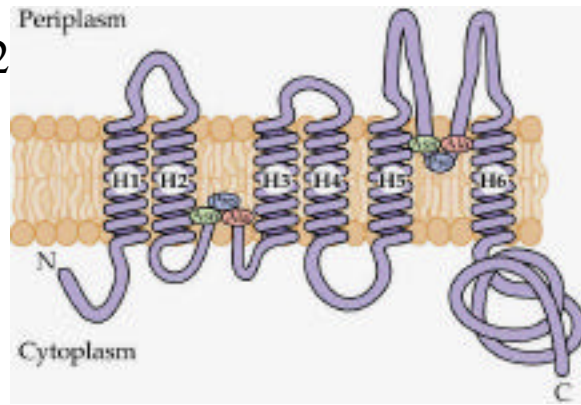
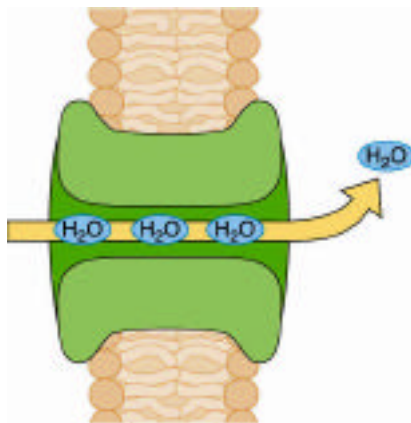


Fig. 3. 50



Loss of turgor

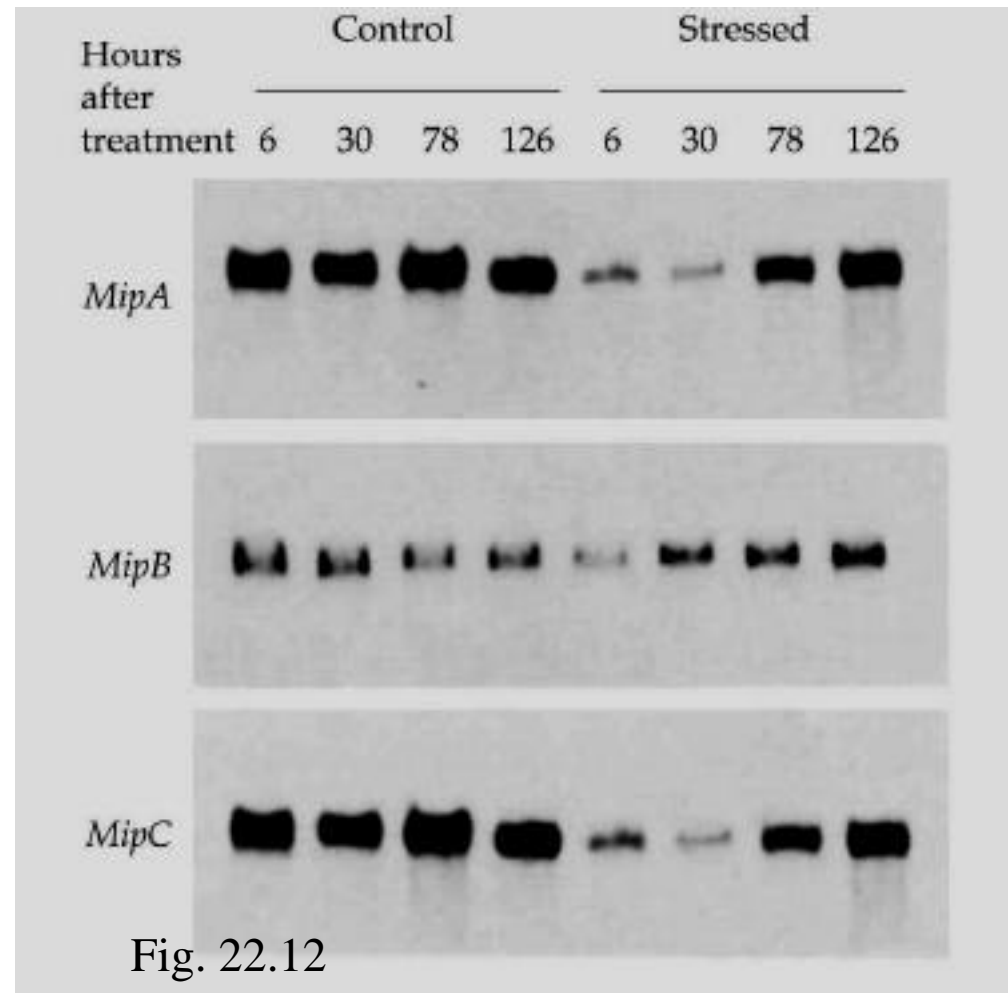
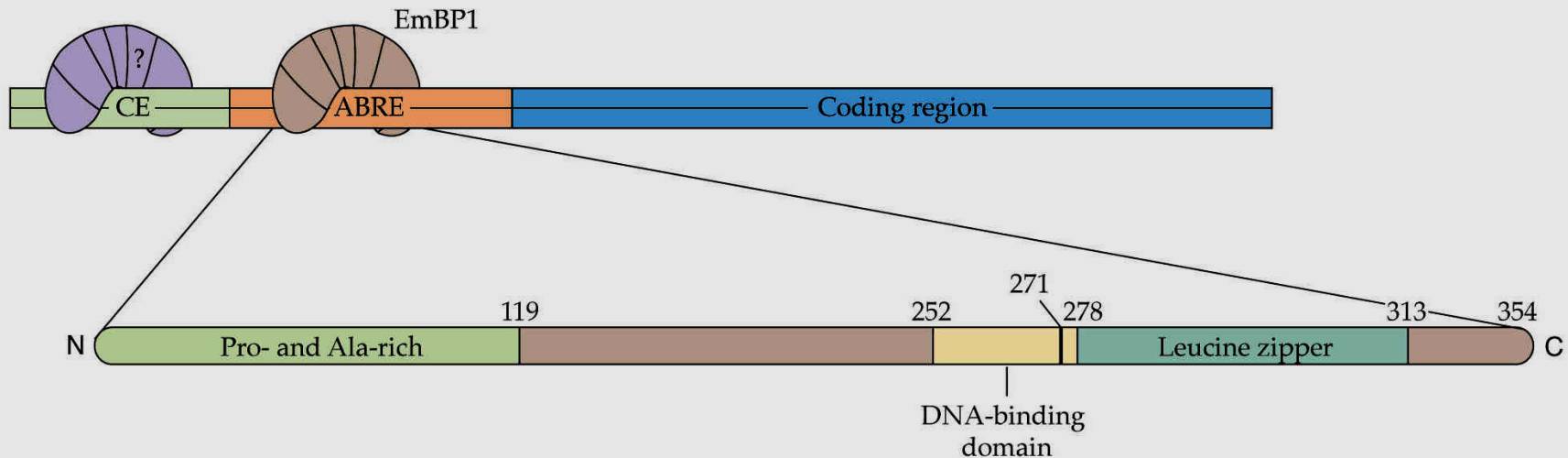


Fig. 22.12

Regulation of MIP expression can regulate permeability of plant cells to water

ABA plays a role in responses to water stress
by regulating stomatal closure and induction of gene expression
but not all water-deficit-responsive genes are induced by ABA

ABA-induced gene



Freezing stress

Chemical potential of ice is less than liquid water

Vapor pressure of extracellular ice is less than that of water in cytoplasm

As ice is initiated in intercellular spaces, water moves out of cytoplasm
toward extracellular space

Freezing tolerant plants:

1. promote extracellular ice formation, preventing ice crystal in cytoplasm
2. Accumulate antifreezing proteins in apoplast, slowing ice formation

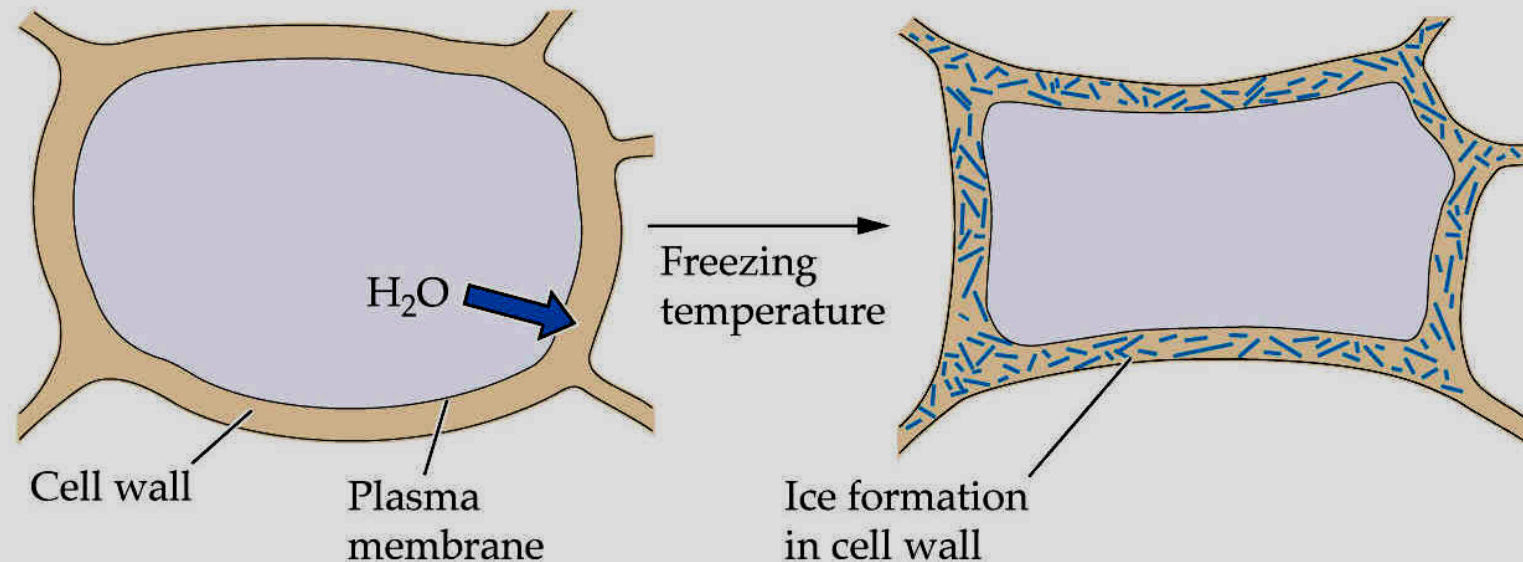


Fig. 22.15

Cold acclimation

1. membrane stabilization:

change in membrane lipid composition,
fatty acid desaturation in membrane phospholipids
change in abundance of membrane sterols and cerebroside

2. osmolytes and antifreeze proteins accumulate

antifreeze proteins form oligomeric complexes, whose large surfaces
interact with ice and inhibit its growth and recrystallization

3. changes in gene expression

still poorly understood

Oxidative stress

Read 1189-1197

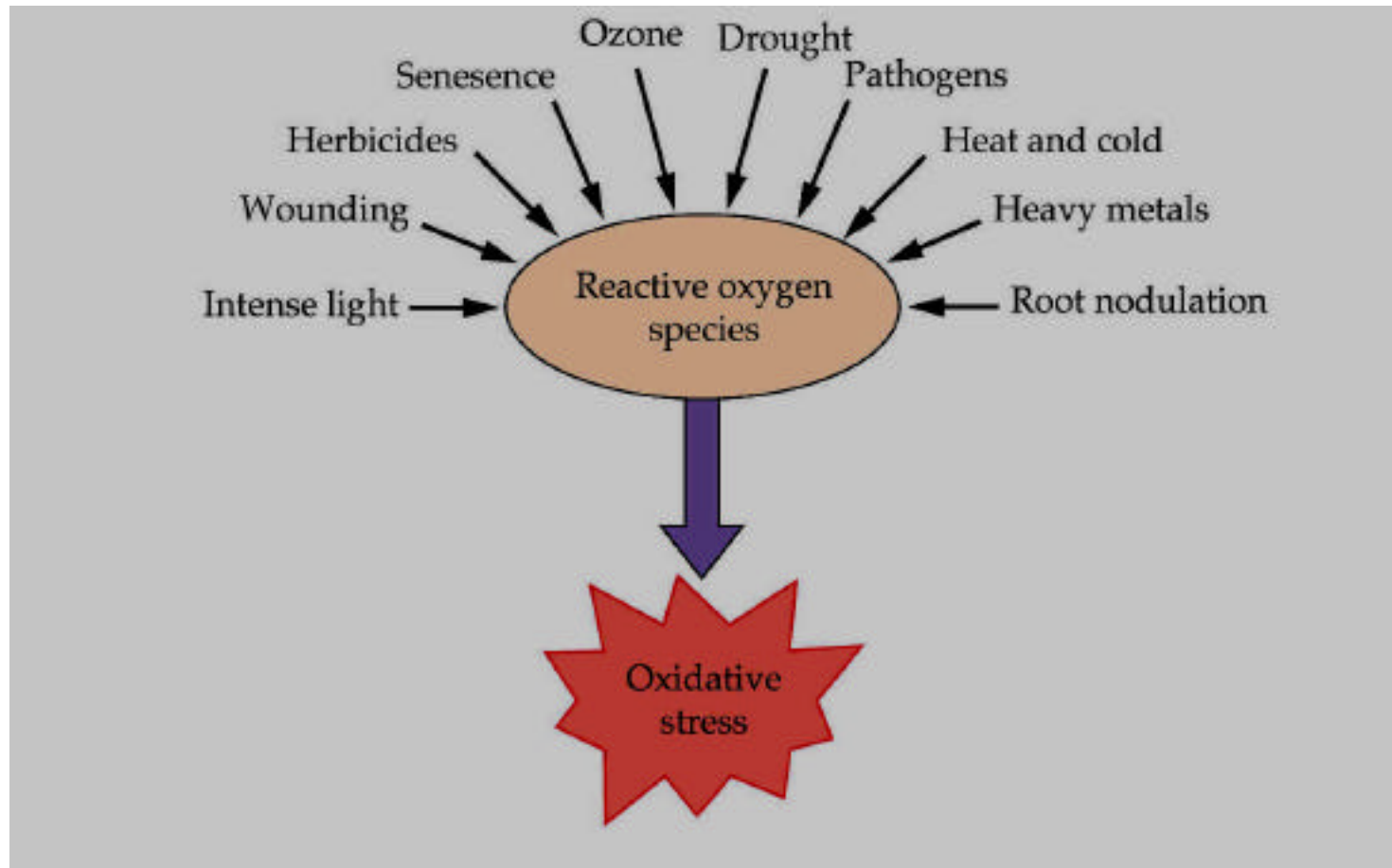
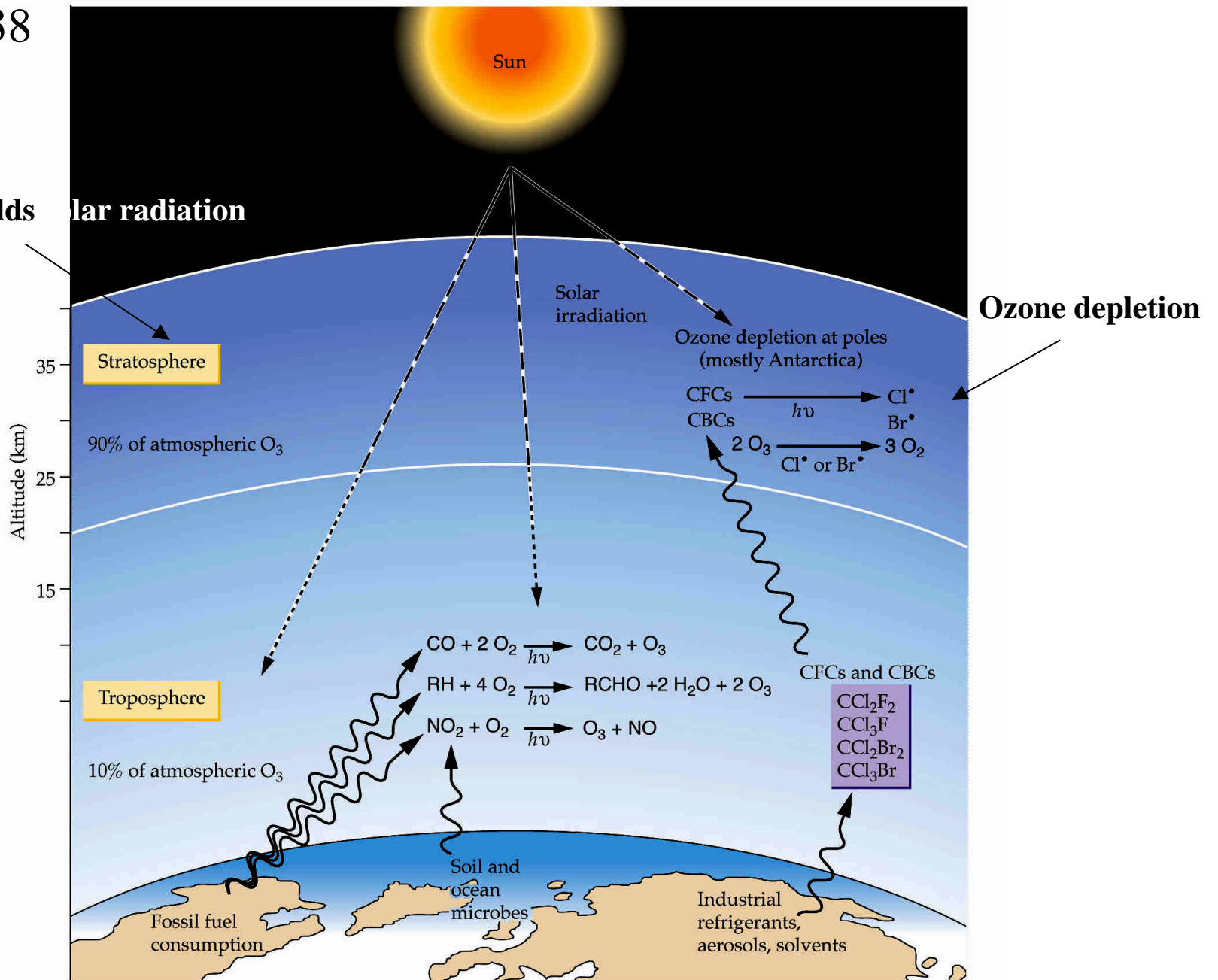


Fig. 22.35

Compound	Shorthand notation(s)	Structural representation(s)	Sources
Molecular oxygen (triplet ground state)	$O_2; {}^3\Sigma$	$\cdot\ddot{O}=\ddot{O}\cdot$ $1s^2 2s^2 (\sigma_s)^2 (\sigma_s^*)^2 (\sigma_x)^2 (\pi_y)^2 (\pi_z)^2 (\pi_y^*)^1 (\pi_z^*)^1$	Most common form of dioxygen gas
Singlet oxygen (first excited singlet state)	${}^1O_2; {}^1\Delta$	$\cdot\ddot{O}=\ddot{O}\cdot$ $1s^2 2s^2 (\sigma_s)^2 (\sigma_s^*)^2 (\sigma_x)^2 (\pi_y)^2 (\pi_z)^2 (\pi_y^*)^2$	UV irradiation, photoinhibition, photosystem II e^- transfer reactions (chloroplasts)
Superoxide anion	O_2^-	$[\cdot\ddot{O}=\ddot{O}\cdot]^-$	Mitochondrial e^- transfer reactions, Mehler reaction in chloroplasts (reduction of O_2 by iron-sulfur center F_x of Photosystem I), glyoxysomal photorespiration, peroxisome activity, plasma membrane, oxidation of paraquat, nitrogen fixation, defense against pathogens, reaction of O_3 and OH^- in apoplastic space
Hydrogen peroxide	H_2O_2	$H-\ddot{O}-\ddot{O}-H$	Photorespiration, β -oxidation, proton-induced decomposition of $O_2^{\cdot-}$, defense against pathogens
Hydroxyl radical	OH^\cdot	$\cdot\ddot{O}-H$	Decomposition of O_3 in presence of protons in apoplastic space, defense against pathogens
Perhydroxyl radical	O_2H^\cdot	$\cdot\ddot{O}=\ddot{O}-H$	Reaction of O_3 and OH^- in apoplastic space
Ozone	O_3	$\begin{array}{c} \cdot\ddot{O} \\ \parallel \\ \ddot{O}^+ \\ \diagup \quad \diagdown \\ \ddot{O} \quad \ddot{O} \end{array}$	Electrical discharge or UV radiation in stratosphere, reactions involving combustion products of fossil fuels and UV radiation in troposphere

Fig. 22.38

Ozone shields solar radiation



Ozone depletion

Altitude (km)

Stratosphere

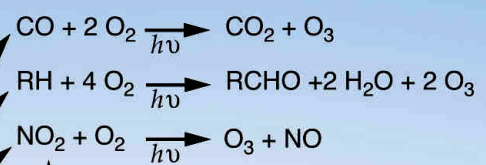
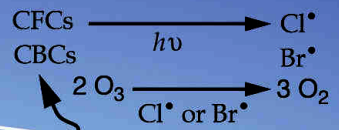
Troposphere

90% of atmospheric O₃

10% of atmospheric O₃

Solar irradiation

Ozone depletion at poles (mostly Antarctica)



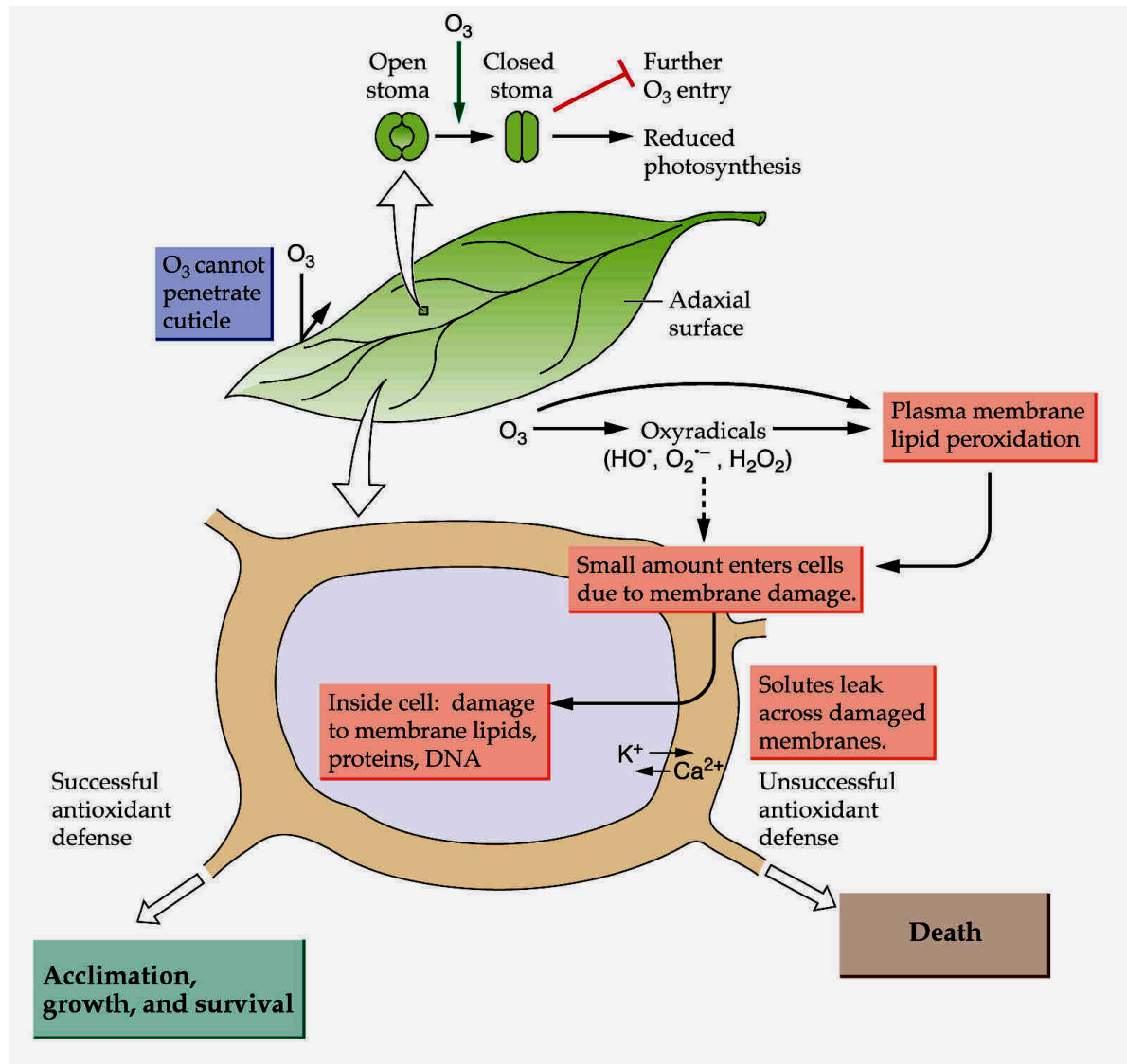
- CFCs and CBCs
- CCl₂F₂
 - CCl₃F
 - CCl₂Br₂
 - CCl₃Br

Fossil fuel consumption

Soil and ocean microbes

Industrial refrigerants, aerosols, solvents

Fig. 22.40



Antioxidants

Ascorbate (Vit. C)

-carotene

Glutathione reduced (GSH)

Polyamine

– Tocopherol (Vit. E)

Zeaxanthin

Subcellular location

Apoplast, cytosol, plastids, vacuole
plastids

cytosol, mitochondrion, plastid

cytosol, mitochondrion, plastid, nucleus
cell membrane, plastid membrane

chloroplast

Antioxidant enzyme:

Ascorbate peroxidase

cytosol, plastid stoma, plastid membrane
root nodule

Catalase

cytosol, glyoxysome, peroxisome

Dehydroascorbatereductase

cytosol, plastid stoma, root nodule

Glutathione reductase

cytosol, mitochondrion, plastid stoma,
root nodule

Monodehydroascorbate
reductase

plastid stoma, root nodule

Superoxide dismutase

cytosol, plastid, peroxisome, root nodule

Table 22.6 and 22.7

Fig. 22.37

