# **RESEARCH PERSPECTIVE**

# Developmental arrest: from sea urchins to seeds

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# Abstract

The phenomenon of dormancy extends beyond the boundaries of the plant kingdom. While plant biologists typically associate dormancy-breaking treatments only with seeds, buds or tubers, these chemicals and environmental stimuli have much broader activity as general terminators of developmental arrest in other, non-plant species. The activation of growth by these treatments is associated with signal transduction processes, metabolic upregulation and changes in gene expression, in addition to other events that may or may not be species specific. The study of both the classic and current developmental arrest literature beyond the boundaries of plant biology may be helpful in generating useful ideas and analogies for meaningful experimental progress towards understanding seed dormancy.

Keywords: developmental arrest, diapause, dormancybreaking chemicals, egg activation, seed dormancy, spore dormancy

# Introduction

To survive adverse environmental conditions (e.g. cold, water limitations, oxygen deprivation, etc.) many species enter an arrested state (e.g. Henis, 1987; Cáceres, 1997 and references therein). Generally, this state can be primarily metabolic arrest alone (quiescence) (Hochachka and Guppy, 1987; Guppy and Withers, 1999) or in combination with

developmental arrest (dormancy or diapause)(e.g. Hand, 1991), where an individual is genetically programmed and specifically prevented from entering subsequent growth stages of its life cycle. Organisms entering developmental arrest generally do so before attaining their adult form. For example, developmental arrest is observed in the egg (Arbacia sp.) (Épel, 1989), blastocyst (*Capreolus* capreolus) (Renfree, 1978), cyst (Artemia sp.) (Drinkwater and Crowe, 1987), larval (Haemonchus contortus) (Petronijevic et al., 1986), gemmule (Eunapius fragilis) (Loomis et al., 1996), pupal (Sarcophaga crassipalpis) (Denlinger et al., 1980), seed (Bewley and Black, 1982, 1994) and spore (Phycomyces blakesleeanus) (Thevelein et al., 1979) stages. However, a developmentally arrested state is not an obligatory component of all life cycles (Sussex, 1978).

Developmental arrest occurs when an organism enters a form resistant to adverse conditions. As growth and development cease, metabolic activities are concertedly down-regulated to minimal levels to retain viability. An ametabolic state does not generally occur, as metabolism can be detected in both dry and hydrated systems (Bewley and Black, 1982; Hand and Gnaiger, 1988; Footitt et al., 1995; Hand and Hardewig, 1996; Brooks and Storey, 1997). Development does not continue when optimum environmental conditions simply resume. An activating stimulus is required to terminate developmental arrest (i.e. to break dormancy). This activating stimulus is usually not required for additional further growth. The nature of developmental arrest has been reviewed in a number of model systems, such as eggs (Loeb, 1913; Epel, 1989, 1990), insects (Jungreis, 1978), seeds (Bewley and Black, 1982; Simpson, 1990; Hilhorst, 1995, 1998; Bewley, 1997), spores (Sussman and Halvorson, 1966) and diapausing aquatic animals (Hand and Podrabsky, 2000).

Therefore, dormancy is a phenomenon common

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from the monera to the animal kingdom. For almost 100 years dormancy research has tended to remain within the bounds of its respective kingdom. This has held back progress in a field that directly and indirectly affects human health and welfare. Progress in our laboratory was contained within the same boundaries until we discovered the sea urchin egg activation research and some early 20th-century seed papers buried in the literature. The conceptual flood gates then opened, and by reviewing the field of dormancy-breaking chemicals without regard to kingdom or species, a pattern emerged that may help to cast light on the underlying mechanism of physiological dormancy-breaking. In the past few years such ideas have started to spread, primarily from the field of egg activation, bringing an increasing awareness of advances in diverse and seemingly unrelated model systems. Focusing upon genetically programmed developmental arrest, this review highlights the widespread nature and commonalities of chemical-based dormancy-breaking treatments and a few consequences of such treatments.

# Early development of chemical-based dormancybreaking treatments

Towards the end of the 19th century, the properties of chemicals capable of perturbing biological systems attracted increasing attention. Fungal spore susceptibility to toxic acids was related to the presence of the undissociated acid form (Clark, 1898). Cell permeability to weak bases was equated with their degree of dissociation and lipophilicity (Harvey, 1911). Alcohol partition coefficients (lipophilicity) and anaesthetic potency were found to increase with carbon number (Overton, 1901). At the same time, developmental biologists recognized these properties as affecting the activity of chemicals that terminated developmental arrest (Loeb, 1913).

# Common thematic features of chemical-based dormancy-breaking treatments

# Structural features

A large number of dormancy-breaking chemicals are active in surprisingly diverse model systems (examples in Table 1). Small molecular weight substances, known to many of us only as seed dormancy-breaking chemicals (e.g. weak acids, aldehydes, ketones and alcohols), terminate developmental arrest in a variety of species. From this survey, the common structural features of dormancybreaking chemicals can be identified (see following sections). In our laboratory, consideration of these features increased both the successful use and discovery of seed dormancy-breaking chemicals. Lipophilicity plots proved especially useful for predicting the concentration range for dormancy-breaking activity of previously untested chemicals (Cohn, 1989, 1997; Cohn *et al.*, 1989, 1991). In addition to lipophilicity, the nature and position of oxidizable functional groups were critical factors governing activity (reviewed in Cohn, 1996a).

# The pH effect and the impact of dissociation constants

Acid scarification, a common seed treatment, terminates developmental arrest in systems other than seeds. A number of reports have shown that inorganic acids at pH 2 break dormancy (Loeb, 1913; Lillie, 1926; Sibilia, 1930; Bingham and Meyer, 1979; Adkins et al., 1985; Petronijevic et al., 1986). Whether activation is a result of acid scarification/mechanical injury or acid loading is not clear. Weak acids and bases also break dormancy under milder, more controlled conditions in a pH-dependent manner, and this effect is related to their dissociation constants (pKs). Dormancy-breaking activity requires the presence of the uncharged chemical species. This has been recognized in a number of model systems (Loeb, 1913; Lillie, 1926; Toole and Cathey, 1961; Palevitch and Thomas, 1976; Cohn and Hughes, 1986; Petronijevic et al., 1986; Van Mulders et al., 1986; Cohn et al., 1987; Petronijevic and Rogers, 1987a).

# Contact time

The application of dormancy-breaking chemicals as a pulse is more effective than continuous contact (Loeb, 1913; Lillie, 1926; Zagorski and Lewak, 1984; Cohn and Hughes, 1986). This is almost certainly due to the detrimental effect of prolonged exposure to dormancy-breaking chemicals on normal development (Loeb, 1913; Lillie, 1926; Mayer and Evenari, 1953). These reports also confirm what has been commonly known for many seeds: conditions for the termination of developmental arrest can be very different from those supporting growth.

### Lipophilicity and molecular size

In several model systems the dormancy-breaking activity of organic acids and their derivatives is correlated with their lipophilicity (Loeb, 1913; Thevelein *et al.*, 1979; Belmans *et al.*, 1983; Taylorson, 1988; Cohn *et al.*, 1989). In seeds, this correlation is modified by a functional group effect, with weak acids being more active (Cohn *et al.*, 1989). The activity of some compounds (e.g. inorganic nitrogen

**Table 1.** Chemicals showing dormancy-breaking activity and the kingdoms in which they are found to be active. In each kingdom, the number designates a species where the appropriate chemical shows activity. The chemicals and species identified are representative only; other examples can be found in the cited literature. Species are identified in the species key below. Within each kingdom, species are arranged by phyla

	Kingdom				
Chemical	Monera	Protista	Fungi	Plantae	Animalia
Alkanes					
Butane	2				
Pentane	2				10
Hexane	2				9, 10
Heptane					10
Iso-octane					10
Cyclohexane					10
Alkenes					
Ethylene				10, 11, 13, 14	
Propylene	2			13, 14	
Propadiene				14	
1-Butene	2				
1-Hexene					10
Aonocarboxylic acids					
Methanoic acid	2		1	7	4, 12, 15
Ethanoic acid	2	2, 3	1,3	1, 3, 7	6, 12, 13, 15
Propanoic acid	2		1, 4	7	12, 15
2-Propenoic acid			1		
Butanoic acid	2		1, 3, 4	1, 2, 7	12, 13, 15
Isobutanoic acid			1, 3	7	12
Isopentanoic acid			1	7	12
Pentanoic acid	2		1, 3	7	12, 15
Hexanoic acid	2		1	7	12, 15
Isohexanoic acid					12
Heptanoic acid					15
Octanoic acid					15
Nonanoic acid					15
Decanoic acid					15
Palmitic acid			9		
Oleic acid			8, 9		
Linoleic acid			8, 9		10
Linolenic acid			8		
Dicarboxylic acids					
Fumaric acid		7			
Malonic		7			
Oxalic acid		7			6, 15
Succinic acid		7		7	
Fricarboxylic acids					
Citric acid		7			15
Hydroxyacids					
Glycolic acid				7	
Ascorbic acid				3, 4	
Lactic acid				7	12
β-Hydroxybutyric acid				7	15
Aldehydes					
Formaldehyde				6	
Acetaldehyde				3,7	
Propionaldehyde			c.	7	
Pentanal			9		
Hexanal			8,9	15	
2,4-Hexadienal			8		
Heptanal			7,9		
Octanal			7, 8, 9	15	
Nonanal			7, 9		
					Continued

Table 1. Continued	Tab	le 1.	Continued
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	Kingdom					
Chemical	Monera	Protista	Fungi	Plantae	Animalia	
Decanal			9			
Hexadecanal			8			
Esters						
Methyl formate				7		
Methyl propionate				7		
Ethyl acetate				7	10	
Ethyl butyrate			2			
Ketones						
Propanone			2	7ª, 9	9, 10	
Butanone				,	10	
2-Pentanone					10	
3-Pentanone					10	
1-Penten-3-one			8, 9			
2-Hexanone			6, 7			
2-Heptanone			6, 7	15		
2-Octanone			5, 6, 7	15, 16		
2-Nonanone			5, 6, 7	15, 16		
Alcohols			, -,	, -		
Methanol	3		1, 2	7, 9, 10, 11		
Ethanol			1, 2	3, 4, 5, 7, 8, 11	8, 12	
Propanol			1, 2	3, 4, 5, 7, 8	12	
Isopropanol			2	7, 11, 16		
2-Propen-1-ol				4		
Butanol	2		1, 2	3, 4, 5, 7	10, 12	
Isobutanol			-, -	7 <sup>a</sup>		
Pentanol	2		1, 2	5, 7, 8		
Hexanol	2 2		1, 2	7 <sup>a</sup>		
Heptanol	-		1, 2	7 <sup>a</sup>		
Octanol			1, 2	7 <sup>a</sup>		
3-Octanol			-) -		1	
1-Octene-3-ol					1	
Nonanol			7		1	
2-Nonanol			7	15, 16		
3-Nonanol			-	16		
1-Nonen-3-ol				16		
Amines				10		
Hydroxylamine	2			3, 7, 10		
Methylamine	1			0,7,10	5	
Ethylamine	2				5	
Propylamine	-			7 <sup>a</sup>	0	
Butylamine				,	5, 13	
Heptylamine	2				0,10	
Octylamine	2 2 2					
Decylamine	2					
Dodecylamine	2					
Ethers	2					
Diethyl ether				9, 11	8, 10, 11	
Aromatics				<i>J</i> , 11	0, 10, 11	
Benzene					10	
Benzoic acid				7	10	
Benzyl acetate			5,7	1	12, 13	
				16		
Benzaldehyde Benzyl alcohol			5,7			
Benzyl alcohol Benzyl amino				5, 8	E 10	
Benzyl amine Phonol			1		5, 13	
Phenol Solignia agid			1	-	10	
Salicylic acid			1	7	12	
Salicylhydroxamic acid Methyl salicylic acid			5, 7	13, 15 10		

# Table 1. Continued

	Kingdom				
Chemical	Monera	Protista	Fungi	Plantae	Animalia
Salicylaldehyde			7		
Toluene					10
Xylene					10
Xylol					8, 11
Inorganics					
Ammonia	1		1	3, 7ª, 12	2, 3, 4, 5, 13
Azide				3, 7, 10	
Carbon dioxide		1, 2, 3, 4, 5, 6		5, 7, 9	2, 3, 4, 7, 12, 13, 14, 15
Cyanide				7, 10, 13	12, 14
Hydrogen peroxide				6	7
Hydrogen sulphide		5		7 <sup>a</sup>	4
Nitrate	2	2, 3		3, 10	12
Nitrite				3, 7, 10	
Nitric oxide		5			
Nitrogen dioxide				7	

<sup>a</sup> Unpublished data.

# Key to species:

### MONERA

Schizophyta: 1, Bacillus cereus endospore (Preston and Douthit, 1988); 2, Bacillus megaterium endospore (Levinson and Sevag, 1953; Rode and Foster, 1961, 1965); 3, Thermoactinomyces vulgaris endospore (Kirillova et al., 1974).

#### PROTISTA

Sarcomastigophora: 1, Giardia muris cyst (Schaefer et al., 1984); 2, Hartmannella rhysodes cyst (Datta, 1979); 3, Schizopyrenus russelli cyst (Datta, 1979).

Apicomplexa: 4, Eimeria bovis cyst (Jensen et al., 1976); 5, Eimeria stiedai cyst (Jensen et al., 1976); 6, Eimeria tenella cyst (Nyberg et al., 1968; Jensen et al., 1976).

Ciliophora: 7, Pleurotricha lanceolata cyst (Jeffries, 1956, 1962).

#### FUNGI

Amastigomycota: 1, *Phycomyces blakesleeanus* conidiospore (Thevelein *et al.*, 1979, 1983; Van Mulders *et al.*, 1986); 2, *Neurospora tetrasperma* ascospore (Belmans *et al.*, 1983); 3, *Hygrophorus russula* basidiospore (Ohta, 1988); 4, *Tricholoma flavovirens* uredospore (French, 1984); 6, *Uromyces vignae* uredospore (French, 1984); 7, *Uromyces rumicis* uredospore (French *et al.*, 1986); 8, *Alternaria alternata* conidiospore (Harman *et al.*, 1980); 9, *Fusarium solani* conidiospore (Harman *et al.*, 1980).

#### PLANTAE

Phaeophycophyta: 1, Fucus vesiculosus egg (Overton, 1913); 2, Sargassum piluliferum egg (Hiroe and Inoh, 1954).

Anthophyta: 3, *Avena fatua* seed (Adkins *et al.*, 1984a, b, 1985; Cairns and De Villiers, 1986); 4, *Avena sativa* seed (Corbineau *et al.*, 1991); 5, *Echinochloa crus-galli* seed (Taylorson, 1988; Leather *et al.*, 1992); 6, *Hordeum vulgare* seed (Hareland and Madson, 1989; Fontaine *et al.*, 1994); 7, *Oryza sativa* seed (Tseng, 1964; Major and Roberts, 1968; Cohn *et al.*, 1983, 1987, 1989; Cohn and Castle, 1984; Cohn and Hughes, 1986); 8, *Panicum capillare* seed (Taylorson, 1989); 9, *Panicum dichotomiflorum* seed (Taylorson and Hendricks, 1979; Taylorson, 1980); 10, *Amaranthus albus* seed (Hendricks and Taylorson, 1974; Taylorson, 1979); 11, *Amaranthus retroflexus* seed (Taylorson, 1979, 1989; Schonbeck and Egley, 1980); 12 *Berbera verna* seed (Hendricks and Taylorson, 1974); 13, *Lactuca sativa* seed (Brooks *et al.*, 1985; Abeles, 1986); 14, *Portulaca oleracea* seed (Taylorson, 1979); 15, *Rumex acetosella* seed (French and Leather, 1979); 16. *Rumex crispus* seed (French and Leather, 1979; Taylorson, 1984; French *et al.*, 1986).

#### ANIMALIA

Nematoda: 1, Bursaphelenchus xylophilus juvenile (Matsumori et al., 1989); 2, Ascaris suum juvenile (Petronijevic and Rogers, 1987a); 3, Haemonchus contortus juvenile (Petronijevic et al., 1986; Petronijevic and Rogers, 1987a, b); 4, Nematospiroides dubius juvenile (Petronijevic et al., 1986).

Annelida: 5. Polynoe sp. egg (Loeb, 1913); 6, Thalassema mellita egg (Loeb, 1913).

Arthropoda: 7, Artenia franciscana cyst (Drinkwater and Crowe, 1987; Clegg et al., 1996); 8, Melanoplus differentialis egg (Slifer, 1946); 9, Manduca sexta pupae (Denlinger et al., 1980); 10, Sarcophaga crassipalpis pupae (Denlinger et al., 1980); 11, Loxostege sticticalis pupae (Pepper, 1937).

Echinodermata: 12, Asterias forbesii egg (Lillie, 1910, 1913, 1926, 1927); 13, Arbacia sp. egg (Lyon, 1903; Loeb, 1913; Harding, 1951); 14, Paracentrotus lividus egg (Lyon, 1903; Loeb, 1913); 15, Strongylocentrotus purpuratus egg (Loeb, 1913).

compounds) appears to be better related to molecular size rather than lipophilicity (Cohn *et al.*, 1989). Overall, it is somewhat disappointing that features, identified by Loeb (1913) in the activation of sea urchin eggs (contact time, dissociation constant and lipophilicity), have taken so long to be recognized in other model systems, especially seeds. On the other hand, detailed structure–activity studies of chemicals that activate eggs have yet to be conducted.

# Is there a common mechanistic basis for crosskingdom chemical activities?

Many plant species exhibit physiological seed dormancy, i.e. resulting not from an impermeable seed coat, physical restraint or an immature embryo, but from the consequences of a developmental programme. Physiological dormancy can be broken in response to environmental, physical and chemical agents (Bewley and Black, 1994). Several hypotheses have been presented to explain the dormancybreaking ability of these agents (reviewed in Bewley and Black, 1982, 1994; Cohn, 1987). In the case of chemical agents, hypotheses tend to centre on specific components of metabolism (e.g. Roberts and Smith, 1977; Taylorson and Hendricks, 1980/81; Esashi et al., 1981a, b). These hypotheses attempt to explain the dormancy-breaking action of specific groups of compounds. However, until recently they have not been able to address the fundamental question of why a wide range of apparently unrelated chemicals are able to break dormancy (see Cohn and Hilhorst, 2000)

Of the known dormant forms in the kingdoms of life, the majority are activated by representatives from many of the chemical classes shown in Table 1. A notable exception has been the alkanes, which appear to have no dormancy-breaking activity in seeds (Taylorson, 1979; Abeles, 1986; Cohn *et al.*, 1989). Initially, it may appear unusual that such a wide variety of chemicals should be active in such a diverse number of species. However, this can be accommodated if similar dormancy-breaking mechanisms operate in all kingdoms.

# Markers for dormancy-breaking and early germination events

In previous publications, we have documented the need for physiological/biochemical markers as guideposts for the time course that seeds experience as they progress from dormancy to germination processes as a consequence of the application of dormancy-breaking chemicals (Footitt and Cohn, 1992, 1995; Cohn and Footitt, 1993; Footitt *et al.*, 1995;

Cohn, 1996a, b). These have included changes in tissue pH, levels of fructose 2,6-bisphosphate, as well as uptake kinetics and metabolism of dormancybreaking chemicals. Review of the historical literature indicates that some of these putative markers have wider relevance beyond the world of seeds.

# Intracellular pH

Many developmentally arrested systems exhibit a change in internal pH upon activation (Table 2). In unicellular systems such as sea urchin and Xenopus eggs, one of the early events of fertilization or artificial activation is an increase in intracellular pH (Grainger et al., 1979; Whitaker and Steinhardt, 1982; Busa and Nuccitelli, 1984; Charbonneau and Grandin, 1989; Epel, 1989; Freeman and Ridgway, 1993; Miller and Epel, 1999). By contrast, intracellular pH decreases upon activation of dormant multicellular systems, such as diapause cysts of the brine shrimp, Artemia franciscana (Drinkwater and Crowe, 1987), larvae and juveniles of the nematodes, Caenorhabditis elegans and Haemonchus contortus (Petronijevic and Rogers, 1987b; Wadsworth and Riddle, 1988), and upon proliferation of Syrian hamster embryo cells (Isfort et al., 1993, 1995). In plants, intracellular pH is also higher in dormant than in non-dormant tissues, e.g. Jerusalem artichoke tuber buds (Gendraud and Lafleuriel, 1983), red rice embryos (Footitt and Cohn, 1992) and barley aleurone cells (Van Beckum et al., 1993). The pH of dormant seed embryos decreased dormancy-breaking treatments during and germination (Table 2). These observations identify changes in intracellular pH as a potential marker for the change in developmental pattern.

#### Small molecular weight substances

At some time interval after the application of a dormancy-breaking chemical to seeds, there must be a resumption of metabolic activity associated with normal germination that occurs prior to radicle emergence. One of the challenges of seed dormancy research is to avoid confusing activities associated with the dormancy-breaking process in contrast to these germination events. Identification of biochemical markers to delineate these processes would be of significant interpretive value. Because of the numerous synthetic processes required for germination, particularly transcription and translation, one of the earliest manifestations of the germination process would be increased carbon flux through energy-generating pathways such as glycolysis (Botha *et al.,* 1992) and an increase in the metabolic regulator, fructose 2,6-bisphosphate (Fru 2,6-P<sub>2</sub>), which activates pyrophosphate:fructose 6phosphate 1-phosphotransferase (PP<sub>i</sub>-PFK) and

Species	pН	Reference		
Unicellular systems				
Strongylocentrotus purpuratus	Î	Johnson <i>et al</i> . (1976)		
Xenopus laevis	Î	Nuccitelli et al. (1981)		
Carcinus maenas	Î	Hervé <i>et al.</i> (1989)		
Urechis caupo	Î	Gould and Stephano (1993)		
Multicellular systems		•		
Crataegus gloriosa	$\Downarrow$	Eckerson (1913)		
Avena fatua	Ų	Atwood (1914)		
Tilia americana	$\downarrow$	Rose (1919)		
Acer saccharinum	Ų	Jones (1920)		
Juniperus virginiana	Ų	Pack (1921)		
Helianthus tuberosa	Ų	Gendraud and Lafleuriel (1983)		
Artemia franciscana	$\Downarrow$	Drinkwater and Crowe (1987)		

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Table 2. Examples of species exhibiting a change in pH upon termination of developmental arrest

inhibits cytosolic fructose 1,6-bisphosphatase (Van Schaftingen, 1987; Stitt, 1990). PP,-PFK-specific activity can increase when oxygen is limiting during seedling growth (Mertens et al., 1990; Mertens, 1991), and this could also occur in the dense tissues of a germinating seed, as seems to be the case in *Phaseolus* vulgaris (Botha and Small, 1987). In dormant tubers and roots, restoration of metabolic activity led to an increase in Fru 2,6-P2 (Van Schaftingen and Hers, 1983; Kowalczyk, 1989). In Phycomyces blakesleeanus spores and seeds of Avena sativa, dormancy-breaking chemical treatments induced a transient rise in Fru 2,6-P<sub>2</sub> (Van Laere et al., 1983; Larondelle et al., 1987). In non-dormant Avena sativa, a transient increase was also seen in the germination phase prior to radicle protrusion (Larondelle et al., 1987). The Avena sativa study used intact caryopses. Thus, the transient increase may result from anaerobic conditions in the embryo resulting from oxygen uptake by microflora or the glumes (Lenoir et al., 1986; Briggs and McGuinness, similar 1992), although under conditions no transient increase was seen in barley (Thornton, Footitt and Bryce, unpublished data). After the application of nitrite or propionaldehyde to break dormancy in red rice grains, a dramatic increase in  $[Fru 2,6-P_2]$  was observed within 4 h of chemical treatment. However, other dormancybreaking chemicals did not elicit increased [Fru 2,6- $P_2$  until the time of radicle protrusion, possibly due to inhibitory side-effects of dormancy-breaking weak acids and their esters on glycolytic activity (Footitt and Cohn, 1995). Considering all of this information together, Fru 2,6-P2 may have potential value as a marker for increased metabolic activity during dormancy-breaking and germination in some wellcharacterized situations. However, the differential

Haemonchus contortus

Oryza sativa

sensitivity of seed tissue  $[Fru 2,6-P_2]$  to various chemical and environmental factors prevents adoption of  $[Fru 2,6-P_2]$  as a 'fail-safe', universal marker for dormancy-breaking or the dormant/non-dormant transition.

Petronijevic and Rogers (1987b)

Footitt and Cohn (1992)

In dry wild oat (Avena fatua) seeds, the adenosine triphosphate (ATP) concentration is lower in dormant than in non-dormant seeds (Adkins and Ross, 1983). On hydration, differences in developmental state were not reflected in the ATP levels or energy charge (Adkins and Ross, 1983; Larondelle et al., 1987; Côme et al., 1988). Gibberellic acid (Adkins and Ross, 1983) and ethanol (Larondelle et al., 1987) broke dormancy, but ATP levels were the same as in dormant seeds. However, in dormant Kalanchoë blossfeldiana seeds, a non-saturating dose of red light increased ATP content without breaking dormancy, in effect inducing a physiological response in a dormant seed (De Greef et al., 1989; Dedonder et al., 1992). As ATP levels are highly buffered, the level of unbound adenosine diphosphate (ADP) has been suggested as a more useful measure of energy status (Guppy et al., 1994). Results to date suggest that ATP content and energy charge of seeds may not be useful as uniform markers for the transition between the dormant and germinating states.

Ammonia production is a recently identified marker for the germination phase that merits further investigation. In non-dormant *Arabidopsis thaliana* seed, ammonia levels increased rapidly 24 h prior to radicle emergence (Garciarrubio *et al.*, 1997; Garciarrubio, personal communication). Further work is required to determine whether such an increase occurs in other species and the timing of such an increase in arrested seeds after a dormancy-breaking treatment.

# Enzyme activity

Enzyme levels and activities might be expected to increase in both the dormant/non-dormant transition and the germination phase. To date, studies on seeds have failed to provide an enzyme as a transitional developmental marker between the dormant and germinating states. This is not the case in other model systems. In tubers, enzyme activities differ between tissue from dormant tubers and tissue slices aged in vitro to induce respiration. In aged tissue, the activity of glycolytic enzymes increased, and their distribution changed from the soluble to the particulate fraction (Moorhead and Plaxton, 1988). In animal systems, studies on the activation of sea urchin and clam eggs have also found changes in enzyme activity and location. Total in vitro activity of glucose 6-phosphate dehydrogenase (G6PD) was the same in extracts from fertilized and unfertilized eggs. However, upon fertilization, its localization changed from the insoluble (inactive) to soluble (active) fraction (Isono, 1963; Isono et al., 1963; Ii and Rebhun, 1982; Swezey and Epel, 1986, 1995). If G6PD was assaved *in situ* using permeabilized eggs, the enzyme activity increased rapidly following fertilization, as did NADPH levels (Epel, 1989; Rees et al., 1996). These egg experiments suggest the need for refinement of the 'grind and find' approach previously used in seed studies. (As a side issue, these egg activation data support an important role of the pentose shunt following fertilization in sea urchin. Therefore, there still may be some life in the pentose shunt hypothesis regarding seed dormancy!).

By identifying a series of markers during and following a dormancy-breaking chemical treatment, a time-line of events can be constructed. The position of events on the time-line will suggest other associated events as markers. Hence, the time-line will also fulfil a predictive function. Ultimately, this approach will make it possible to unravel the events that make up the dormancy-breaking process and identify the transition points between the developmental states.

### Overview

While there is a tendency to view dormancy-breaking events as a single linear time course, it now seems that multiple, rapidly interacting paths with different activation kinetics are triggered in response to a single dormancy-breaking signal (Footitt and Cohn, 1992, 1995; Footitt *et al.*, 1995; Cohn, 1996a) and during the germination process as well (Alvarado *et al.*, 2000; Bradford *et al.*, 2000). If constitutive metabolism is activated rapidly above the cellular maintenance level after application of a dormancy-breaking chemical, it is intriguing to ponder what regulatory changes are occurring at the level of transcription and translation (including post-translation) to the control of gene expression during the transition from the dormant to non-dormant state and subsequent germination (Morris *et al.*, 1991; Goldmark *et al.*, 1992; Hance and Bevington, 1992; Dyer, 1993; Johnson *et al.*, 1995; Li and Foley, 1995, 1996; Aalen, 1999; Holdsworth *et al.*, 1999; Alvarado *et al.*, 2000; Bradford *et al.*, 2000).

In addition, one must further consider the role of abscisic acid (ABA), produced in the embryo, which is required to induce dormancy (e.g. LePage-Degivry and Garello, 1992; Hilhorst, 1995; McCarty, 1995; Koornneef et al., 1998). ABA induces increased cellular pH (Gehring et al., 1990; Van der Veen et al., 1992) and the expression of dormancy-associated genes (Morris et al., 1991; Goldmark et al., 1992; Li and Foley, 1995), including the gene for the transcription factor VP1 and its homologues which control embryo maturation and, in some cases, developmental arrest (McCarty, 1995; Jones et al., 1997). This suggests the possible involvement of increased embryo pH in the expression of genes responsible for the induction/ maintenance of seed dormancy. In this respect, it is intriguing that dormancy-breaking weak acids acidify the internal pH (Cohn et al., 1989; Footitt and Cohn, 1992) and inhibit ABA-induced gene expression (Van der Veen et al., 1992). In contrast, gibberellic acidinduced gene expression did not appear to be pH sensitive (Van der Veen et al., 1992; Heimovaara-Dijkstra *et al.*, 1995). These data suggest that internal pH may act as a modulator of some developmental signals, as in the proliferation of Syrian hamster embryo cells, where transient intracellular acidification is indispensable for platelet-derived growth factor-induced proliferation (Isfort et al., 1995).

The role of cellular acidification in the loss of dormancy may be to reduce the expression of ABArelated genes, e.g. VP1 and PKABA1 homologues. In dormant wild oats, transcription of the *afVP*1 homologue is reduced as a result of dry after-ripening and increased on induction of secondary dormancy (Jones et al., 1997, 2000). VP1, as well as PKABA1, represses transcription of germination-specific amylase genes (Hoecker et al., 1995; Walker-Simmons, 1998, 2000; Gómez-Cadenas et al., 1999). VP1 has also been implicated in repression of key glyoxylate cycle genes (isocitrate lyase and malate synthase) during seed development, thus restricting reserve lipid breakdown prior to germination (Paek et al., 1998). Thus, it is possible that VP1, PKABA1 and their homologues have a wider role in repressing the expression of germination-specific genes. The effect of dormancy-breaking chemicals may be to downregulate expression of VP1 gene homologues or to interfere with the activity of the transacting factor itself, via cellular acidification. The slow rate of

germination observed following some dormancybreaking treatments (Footitt and Cohn, 1992, 1995; Myers *et al.*, 1997) may result from the effects of residual VP1 homologue gene products, especially in highly dormant lines (Jones *et al.*, 1997), and the secondary toxicological effects of dormancy-breaking chemicals.

To conclude, we have begun to identify potential molecular, biochemical and physiological markers that can serve as orientation points for further dissection of the dormancy-breaking process, the dormant/non-dormant transition, and germination processes prior to radicle emergence in seeds. While most of the markers identified to date are components of the germination process, some progress has been made concerning the progression of seeds from the dormant to the non-dormant state. Based upon parallels to the termination of development arrest in a wide range of organisms in addition to plants, it is highly likely that general components of the current signalling paradigm (calcium, G proteins, nitric oxide, protein kinases and phosphatases, etc.) will play a significant role, as already described extensively for the aleurone model system (Ritchie et al., 2000). The results of the literature survey presented here suggest a role for seed cytosolic pH changes. For future seed research, it will be important to confirm experimentally that existing markers are widely relevant to a range of species with various dormancy syndromes, as would be predicted from our crosskingdom review of activating chemicals. As dormancy appears to be so similar in many embryonic systems, it is also intriguing to wonder whether homologues to genes such as VP1 and PKABA1 fulfil related functions in Artemia cysts, nematodes, unfertilized oocytes and diapausing insects.

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