

Exploring the physiological basis of cryopreservation success and failure in clonally propagated *in vitro* crop plant germplasm

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An appraisal of potato and *Ribes* shoot meristem cryopreservation shows physiological factors influence survival and development, sometimes independently of protocol and genotype. Markers for oxidative damage incurred by cryostorage reveal two responses: (1) oxidative stress with an eventual decline in regrowth and (2) an oxidative burst associated with higher survival. Differential responses to cryoinjury are discussed in relation to *in vitro* ageing and genetic stability within the conceptual framework of cryobionomics. The possibility that cryopreservation-induced cell death and apoptosis occurs in plants is considered with respect to current concepts of animal cell cryoinjury. It is proposed that a more holistic approach is now required to understand the basis for success or failure of cryopreserved plant germplasm.

Key-words: apoptosis, ageing, clonal, cryopreservation, physiology, potato, *Ribes*, stability

Introduction

Cryostorage success cannot always be attributed to genotypic or cryogenic factors (cryoprotection, cooling and freezing) therefore, it is vital to understand the role of decisive physiological factors

in cryopreservation failure. Several reasons may account for this: (1) restricted application to highly intolerant genotypes within a crop; (2) variable responses within and across genotypes; (3) low number of samples surviving cryostorage; (4) inability of meristems to regrow shoots and (5) idiosyncratic variability within and between experiments. While

an acceptable number of germplasm accessions held in working cryobanks respond satisfactorily to cryopreservation, the application of otherwise successful protocols to some genotypes results in low survival (Keller et al. 2008, Panis et al. 2005). This has proven to be a persistent problem across different species. To address these issues, we examine the responses of clonal crop germplasm within the remit of the European CRYOPLANET - COST Action 871 project (Panis 2007). The effects of cryogenic and non-cryogenic factors are considered specifically for the recovery of potato and *Ribes*, as they provide exemplars of tuber and woody perennial crops. Shoot tip meristem responses to cryopreservation are evaluated using a reappraisal of the authors' earlier potato literature in the context of new studies on *Ribes*.

Materials and Methods

In vitro shoot cultures of *Solanum tuberosum* cultivars (cvs) 'Golden Wonder' and 'Desiree' were established on Murashige and Skoog (MS) medium (Benson et al. 1989). The effects of pre- and post-cryopreservation light (high/low irradiance) treatments were studied on cryoprotected meristems, which were cryopreserved by ultra-rapid freezing (Benson et al. 1989) or controlled rate cooling (Harding et al. 1991) followed by thawing and recovery on MS media containing different hormone combinations (Harding and Benson 1994). Recovery of potato plants as time courses for *in vitro* regrowth and maturation were constructed as reported by Harding and Benson (1994) and Harding (1997). Apical shoot cultures of wild *Solanum* species: *S. phureja*, *S. brachycarpum*, *S. acaule*, *S. guerreroense*, *S. iopetalum* and *S. tuberosum* cv 'Pentland Squire' were cryopreserved using alginate encapsulation-dehydration (Benson et al. 1996). Genotypes of *Ribes ciliatum* (sensitive) and *R. nigrum* cv 'Ben More' (tolerant) were selected based on their differential responses to cryopreservation (Reed et al. 2005). *In vitro* shoots were grown on MS-*Ribes* medium and shoot meristems sucrose-acclimated

(Johnston et al. 2007) and cryopreserved using encapsulation-dehydration (Reed et al. 2005). Stress marker profiles were constructed using antioxidants and volatile hydrocarbon analyses of methane (a marker for •OH) and ethylene (Johnston et al. 2007). Profiles were constructed during sucrose-simulated acclimation and after cryostorage.

Results

Potato genotypic effects

Differential genotype responses were observed in *S. tuberosum* cvs 'Golden Wonder' and 'Desiree', with 'Golden Wonder' being consistently more tolerant to cryopreservation irrespective of light or cryogenic treatments (Fig. 1). Both genotypes exhibited a progressive light-independent decline during recovery, with an onset at week 3 for cvs 'Golden Wonder', whereas 'Desiree' had consistently low levels of recovery across all light conditions (Fig. 1A). Light treatments had a significant effect on total recovery (mean % values) of both cvs 'Desiree' ($p=0.001$, SEM=1) and 'Golden Wonder' ($p=0.025$, SEM=2.9). A very significant interaction was observed (Fig. 1B) between freezing method, post-light treatment and time of total recovery (mean % values) of cv 'Golden Wonder' ($p=0.001$, SEM=4). Overall, ultra-rapid freezing supported higher levels of recovery across both genotypes and high light triggered a decline in recovery and regrowth from week 3 onwards.

Effects of *in vitro* age

Age of culture (Fig. 2A) had a very significant effect on total recovery (mean % values) in both cvs "Desiree" ($p=0.001$) and 'Golden Wonder' ($p=0.001$). Higher survival was observed in younger (short-term) cultures cryopreserved within 6–8 weeks of their initiation in culture from tuber sprouts. Survival was reduced in older (long-term)

cultures maintained for 3 years. Young cultures of both cultivars performed better with respect to shoot production in surviving meristems (Fig. 2B). Culture age had a very significant effect ($p=0.001$) on

cv ‘Golden Wonder’ for which meristem regrowth from younger cultures (Fig. 2B) occurred more rapidly and with a higher % regeneration compared to older cultures.

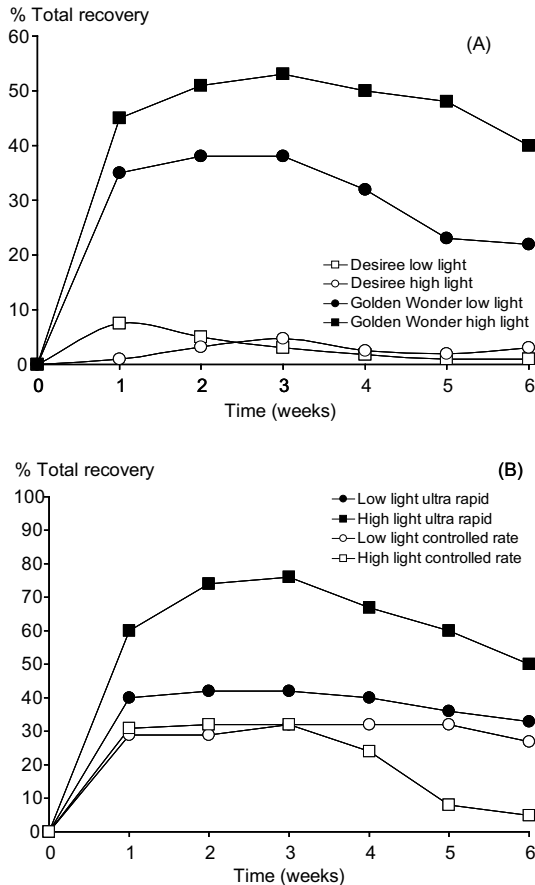


Fig. 1. Time course of (A) the effects of post-light on total recovery of cryopreserved shoot tips of *S. tuberosum* cvs ‘Desiree’ and ‘Golden Wonder’ and (B) the interaction between freezing method, post-light and time on total recovery of *S. tuberosum* cv ‘Golden Wonder’. Data represented as mean (%) values, probability (p) and standard error of the mean (SEM) of three replicated experiments each of 25 shoot tips per light treatment. Low light treatment refers to photosynthetically active radiation (PAR) $15 \mu\text{E}\cdot\text{m}^{-2}\cdot\text{sec}^{-1}$ and high light to PAR $45 \mu\text{E}\cdot\text{m}^{-2}\cdot\text{sec}^{-1}$ original data from Benson et al. (1989) with permission of CryoLetters.

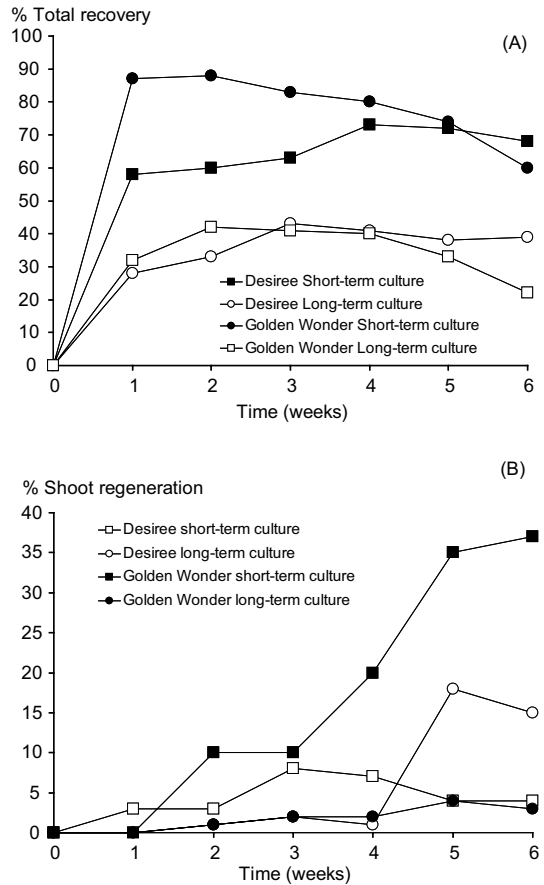


Fig. 2. The effect of pre-freeze short- (6–8 weeks old *in vitro* cultures) and long-term (~3 year old *in vitro* cultures) culture time on (A) total recovery of cryopreserved shoot tips of *S. tuberosum* cvs ‘Desiree’ and ‘Golden Wonder’ and (B) shoot regeneration of *S. tuberosum* cvs ‘Desiree’ and ‘Golden Wonder’. Data represented as mean (%) values and probability (p) of 20 replicate plates derived from 100 shoot tips per treatment original data from Harding et al. (1991) with permission of CryoLetters.

Experimental variability

Significant variation between triplicate experiments and replicate culture plates within an experiment were observed for cv ‘Golden Wonder’ shoot tips cryopreserved using ultra rapid freezing (Fig. 3). In sequence, variation (mean % values) was observed for total recovery ($p=0.001$, SEM=3.2), shoot regeneration ($p=0.001$, SEM=1.4) following 6 weeks of culture and plantlet formation ($p=0.05$, SEM=1.6) over 12 months. Encapsulation-dehydration of shoot tips excised from *in vitro* cultures of wild *Solanum* species resulted in wide variation between triplicate experiments and replicates (mean % values) with respect to total recovery and shoot regeneration after cryopreservation (Fig. 4).

Dynamics of recovery and molecular analysis of potato

Growth regulator composition affected growth and development during recovery of cv ‘Golden Wonder’ shoot tips after cryopreservation (Fig. 5A–C). TBZ medium proved to be the most successful supporting a more synchronized and uniform pattern of recovery. The sequential stages of recovery as mean times for: (i) *in vitro* growth (ii) plant maturity and (iii) total time to regenerate and mature with mean (%) regeneration data for cv ‘Golden Wonder’ are shown in Table 1. Data ranked in order of increasing total recovery time show approximately a 100 day difference in recovery between media containing different hormones. Differences in DNA extracted from mature *S. tuberosum* plants are presented in Figure 5D as an autoradiograph of a Southern blot of the ribosomal RNA gene (rDNA) fragments detected using a 4.4kb ribosomal probe derived from pTa 71. DNA samples (A–M) correlate respectively with plants recovered from cryopreserved shoot tips of cvs ‘Golden Wonder’ and ‘Desiree’ that had been recovered on different hormone regimes. The inset (arrow) marks the variable 2.55kb rDNA fragment in a number of samples (B, D, F, I, J and L) and their respective densities are shown in Table 2. A reduction in the 2.55kb fragment density was

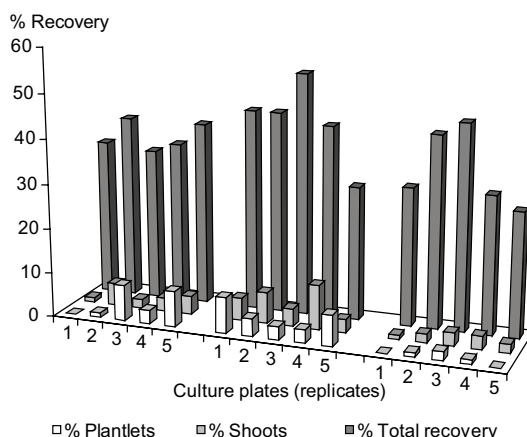


Fig. 3. Variation in total recovery, shoot regeneration and plantlet formation between triplicate experiments and replicate culture plates within an experiment of cryopreserved shoot tips of *S. tuberosum* cv ‘Golden Wonder’. Data represented as mean (%) values, probability (p) and standard error of the mean (SEM) of three replicated experiments each of 25 shoot tips original data from Benson et al. (1989) with permission of CryoLetters.

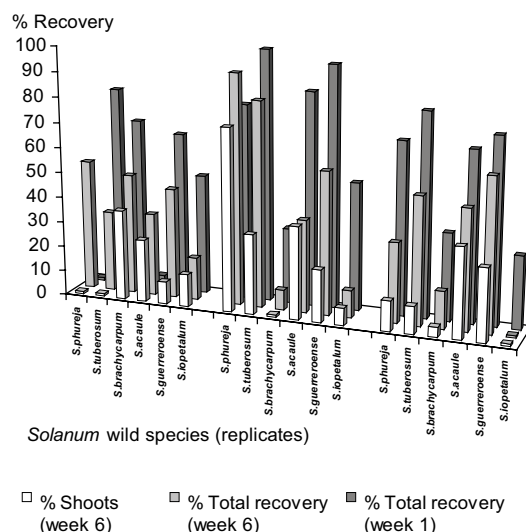


Fig. 4. Variation in total recovery and shoot regeneration between triplicate experiments and replicate samples within an experiment of cryopreserved shoot tips of diverse wild *Solanum* species. Data derived from three replicated experiments, where $n = 25-40$ shoot tips per replicate original data from Benson et al. (1996) with permission of CryoLetters.

observed in samples B, D, F, I, J and L (shown in bold) corresponding to >2.00 ratio for the other (3.1, 1.7 and 0.65kb) rDNA fragments. The respective positions for recovery of samples E, F, I and

J during *in vitro* culture (A), their maturation (B) and total time to regenerate and mature (C) are marked in Figure 5.

Table 1. *In vitro* culture time, maturation time of plantlets, total time of plants to (%) regenerate and mature following cryopreservation of *S. tuberosum* cvs Golden Wonder and Desiree shoot tips recovered on media containing different hormone regimes.

Media	<i>In vitro</i> time	Maturation time	Total time	Regeneration (%)
Golden Wonder				
TBZ	94	108	202	42.7
GA3	132	108	240	12.0
HF-BSA	142	109	252	13.3
NAA/GA3	159	109	269	12.0
GA3-BSA	132	170	302	2.7
HF	177	126	303	17.3
GA3/IAA	189	125	314	2.7
Desiree				
GA3/IAA	118	104	222	2.0
HF-BSA	126	97	223	0.7
GA3	132	98	231	2.7
TBZ	164	120	284	2.7
NAA/GA3	169	117	287	3.3
HF	202	101	304	1.3
GA3-BSA	164	154	318	1.3

Table 2. Ribosomal RNA gene fragment densities/ratios for cryopreserved *S. tuberosum* samples

Fragment size (kb)	Samples (density of fragments)												
	A	B	C	D	E	F	G	H	I	J	K	L	M
3.1	0.70	0.78	0.88	0.66	0.94	0.84	0.93	1.07	0.87	0.86	0.95	0.84	0.91
2.55	0.52	0.35	0.84	0.33	0.69	0.17	0.77	0.82	0.26	0.30	0.72	0.13	0.70
1.70	0.65	0.70	0.82	0.57	0.72	0.54	0.66	0.68	0.66	0.62	0.67	0.68	0.75
0.65	0.86	0.91	1.02	0.89	0.94	0.90	0.94	1.14	0.96	0.90	0.98	0.92	0.98
	Samples (ratios)												
3.1	1.35	2.23	1.05	2.00	1.36	4.90	1.21	1.31	3.30	2.90	1.31	6.46	1.30
2.55													
1.70	1.25	2.00	0.98	1.72	1.04	3.20	0.86	0.83	2.51	2.10	0.93	5.23	1.07
0.65	1.65	2.60	1.21	2.70	1.36	5.30	1.22	1.39	3.96	3.00	1.36	7.08	1.40

Original data from Harding (1997) re-used with CryoLetter's permission.

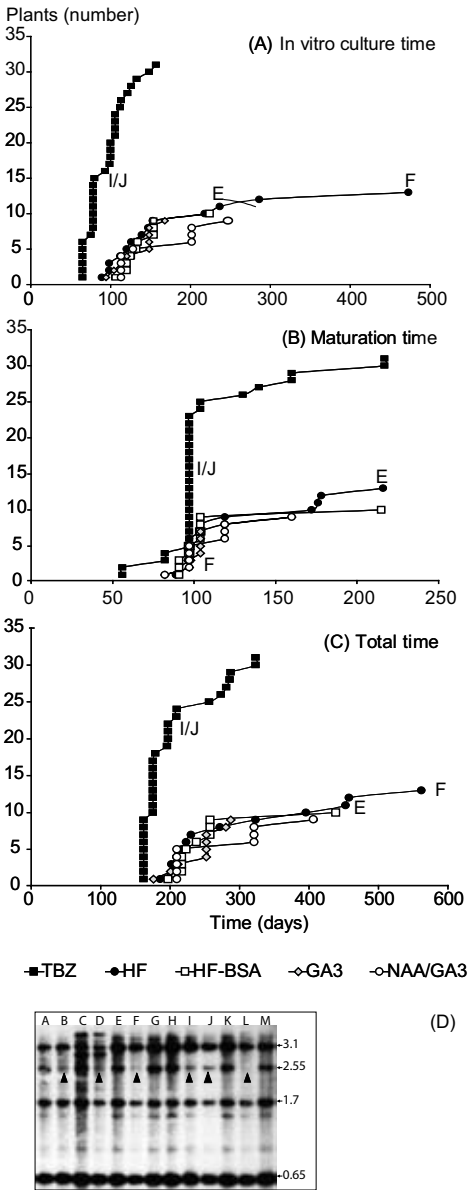


Fig. 5. *In vitro* culture time of plantlets (A), maturation time of plantlets (B) and total time of plants to regenerate and mature (C) and ribosomal RNA gene fragments detected in *S. tuberosum* plants (A–M) recovered on media containing different hormone regimes (D) following cryopreservation. Data represented as mean (%) values of three replicated experiments each of 25 shoot tips per treatment for recovery media: HF, HF-BSA, GA3, GA3/NAA and TBZ original data from Harding and Benson (1994) and Harding (1997) with permission of CryoLetters.

Recovery and oxidative stress in *Ribes* genotypes

The recovery of sucrose-acclimated *Ribes* shoot tips demonstrates *R. nigrum* is more tolerant to cryopreservation, whereas the time course shows a progressive decline in total recovery for the *R. ciliatum* genotype (Fig. 6). Following exposure to successive stages of the encapsulation-dehydration protocol indicates variable stress responses with differential shoot regrowth between the sensitive (*R. ciliatum*) and tolerant (*R. nigrum*) genotypes (Fig. 7 inset). Markers for oxidative stress and antioxidant status (Table 3) revealed significant differences between the two genotypes which can be delineated into two stress profiles (Fig. 7).

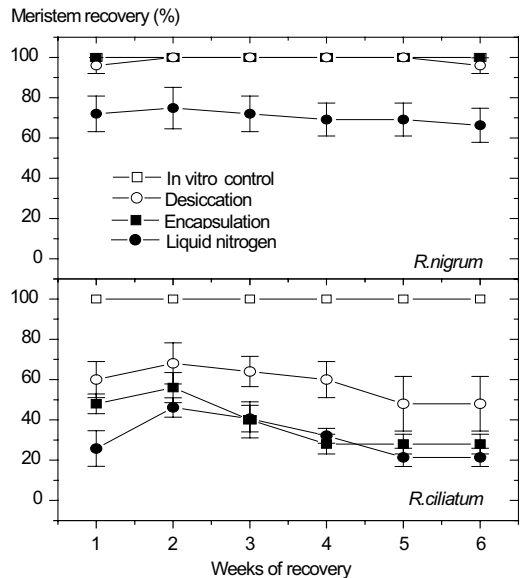


Fig. 6. Time course of cryopreservation tolerant (*R. nigrum* cv Ben More) and sensitive (*R. ciliatum*) shoot tips recovering from different stages of encapsulation-dehydration following 1–2 weeks (as unbleached shoots), 3–6 weeks (as green meristems and expanded leaves) of culture. Values (%) are means and standard errors of the means, where $n = 5$ replicates with 5–8 meristems per replicate original data from Johnston et al. (2007) with permission of Elsevier.

Table 3. Oxidative stress markers and antioxidant status in *Ribes* genotypes following acclimation and cryopreservation.

Stress markers & antioxidants	<i>R. ciliatum</i> (sensitive)		<i>R. nigrum</i> (tolerant)	
	Acclimation	LN	Acclimation	LN
Chlorophyll a ¹	47.6 ± 2.3	-	67.1 ± 5.6	-
Chlorophyll b ¹	25.5 ± 1.0	-	45.4 ± 3.7	-
Chlorophyll a:b ratio	1.9 ± 0.1	-	1.5 ± 0.1	-
Carotenoids ¹	21.2 ± 2.1	-	31.1 ± 2.4	-
Methane ³	11.4 ± 1.5	11.0 ± 5.8	33.6 ± 9.0	12.9 ± 5.1
Ethylene ³	4.5 ± 0.7	42.5 ± 6.8	6.2 ± 3.1	14.6 ± 4.7
Total protein + PVPP ¹	3.83 ± 0.4	4.8 ± 0.2	1.1 ± 0.2	0.5 ± 0.4
Total protein - PVPP ¹	1.9 ± 0.2	2.6 ± 0.7	0.7 ± 0.1	1.8 ± 0.5
Non-phenolic antioxidants ²	5.6 ± 0.3	2.4 ± 0.1	11.7 ± 0.3	4.6 ± 0.7
Total antioxidants ²	29.8 ± 2.2	6.5 ± 1.1	101.6 ± 7.3	65.9 ± 10.9
Total phenolics ¹	19.6 ± 0.4	5.8 ± 0.5	52.4 ± 2.3	53.3 ± 8.1
Protein SH ²	0.2 ± 0.1	1.7 ± 0.1	0.4 ± 0.2	2.5 ± 0.1
Non-Protein SH ²	0.8 ± 0.1	0.2 ± 0.1	1.6 ± 0.1	0.4 ± 0.1
Total SH ²	0.9 ± 0.1	2.2 ± 0.3	1.9 ± 0.2	2.9 ± 0.1

Summary data of oxidative and photooxidative (as pigments) stress markers, proteins and antioxidant levels (units: 1 - mg.g FW⁻¹; 2- μmol.g FW⁻¹; 3 - pmol.s.kg⁻¹) of *Ribes* sensitive and tolerant genotypes following sucrose-simulated cold acclimation (7 days, 0.75M Suc-RIB medium) and cryopreservation (LN) using cryoprotective encapsulation-dehydration (PVPP = polyvinylpyrrolidone, SH = sulphhydryl groups). Original data from Johnston et al. (2007) kindly re-used with Elsevier's permission.

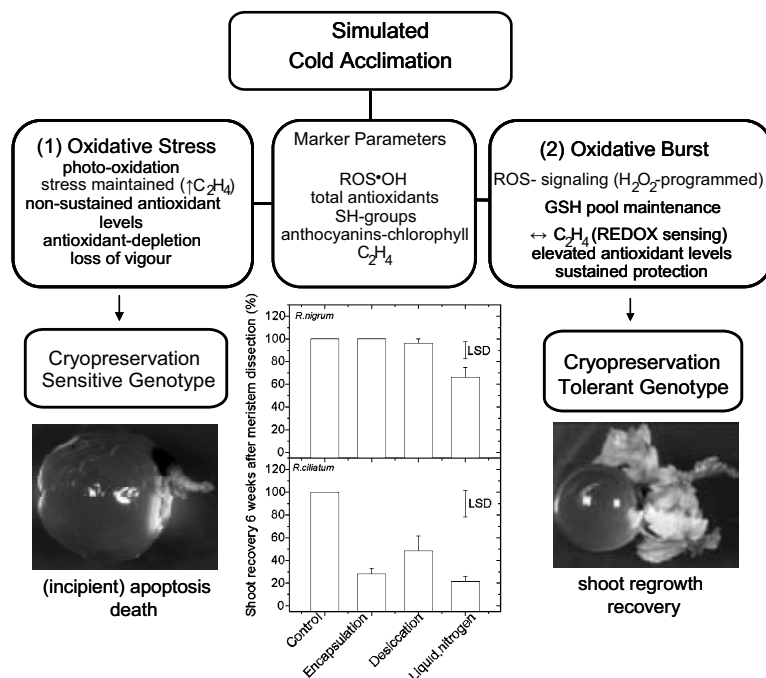


Fig. 7. Model of *Ribes* responses to cryopreservation based on oxidative stress and antioxidant markers. Insert graphic: shoot production of *R. nigrum* (cv Ben More) and *R. ciliatum* meristems recovered on RIB medium, as a percentage (%) of meristems regenerating morphologically normal shoots 6 weeks following exposure to different stages of cryopreservation. Values are means (n = 5 replicates, 5–8 meristems per replicate), standard errors of the mean and 5% (LSD) least significant differences. *P* values for stages of the protocol *R. nigrum* (*p* = <0.001) and *R. ciliatum* (*p* = <0.001) original data from Johnston et al. (2007) with permission of Elsevier.

Discussion

Significant progress in protocol development has resulted in cryopreservation being more widely adopted in genebanks (Kim et al. 2006, Reed 2008) however, for some germplasm variable responses and recalcitrance remain limiting factors affecting cryobank efficiency. Some variability is attributed to known operational factors (Reed et al. 2001, 2004, 2005) and minor, often unpublished technical details (Keller et al. 2008). This comparative study of *Solanum* and *Ribes* also suggests that genotype, protocol, culture regime (light and growth regulator), age of culture and intrinsic experimental variation influence meristem survival and development following cryopreservation. This concurs with previous reports (Schäfer-Menuhr et al. 1997, Golmirzaie and Panta 2000, Keller 2005, Reed et al. 2005).

Predetermined and genotype factors

The droplet (PVS2) vitrification protocol (Panis et al. 2005, Kim et al. 2006) is proving widely applicable across a diverse range of crops, genotypes and cultivars (Sakai and Engelmann 2007). However, variable (intra/inter-species) genotypic responses to cryopreservation can limit its use to certain crops for which widely applicable, validated protocols are desirable. Successful protocols have been largely developed by empirical means however, proteomics now reveals tolerance to cryopreservation has a molecular-physiological basis (Carpentier et al. 2007). Where, genotype is a determinant in recalcitrant germplasm the use of suboptimal protocols may exacerbate and contribute to cryopreservation failure. Comparisons of recovery in *S. tuberosum* cvs 'Desiree' and 'Golden Wonder' (Figs. 1, 2) consistently show 'Desiree' is less amenable to cryopreservation across treatments. Similarly, other studies of cv 'Desiree' demonstrate different survival/shoot (%) responses including: controlled rate freezing ~2/0% (Benson et al. 1989), ultra-rapid freezing ~55/10% (Harding et al. 1991), droplet-freezing ~41/25% (Schäfer-Menuhr et al. 1997), 33/21.7%

(Kryszczuk et al. 2006) and 10–21.4/0% (Keller and Dreiling 2003), encapsulation-dehydration 78.8/58.8% (Grospietsch et al. 1999) and vitrification 70/38.4% (Kryszczuk et al. 2006). Although, protocol refinement enhanced survival this did not necessarily convert to increased development of shoots and plantlets in cv 'Desiree'. This effect was also observed in wild species of potato following shoot cryopreservation using encapsulation-dehydration (Fig. 4) suggesting it may be due to intrinsic stress factors. This is supported by Martinez et al. (2001) who conducted a comparative study on frost sensitive *S. tuberosum* and frost resistant *S. curtilobum* showing higher antioxidant activity (as superoxide dismutase) in the resistant native Andean potato species to freezing and drought. Studies of frost tolerant and intolerant Norway spruce (*Picea abies*) also revealed pre-existing high levels of antioxidants in seedlings contributed to enhanced tolerance in the frost-resistant genotype (Blödner et al. 2005). Cultivars of black currants are distinguished for breeding purposes by their differential sensitivities to spring frosts (Keep et al. 1983) and this helps the selection for intrinsic frost resistance genetic traits in the crop (Mather et al. 1980). The cryopreservation tolerant *R. nigrum* cv 'Ben More' used in this study (Figs. 6, 7) has been previously tested as flower bud frost resistant by Dale and Heiberg (1984). Comparisons of oxidative stress and antioxidants of *R. ciliatum* and *R. nigrum* ('Ben More') similarly suggest differential responses are genetically predetermined. Furthermore, wild and cultivated *Ribes* genotypes display variable recovery responses ranging from zero survival to 90% shoot regrowth following exposure of meristems to liquid nitrogen (Reed et al. 2001, 2004, 2005).

In Vitro Physiology

It is vitally important to recognise that cryopreservation success falls into two distinct phases: (1) the initial shoot tip survival, i.e. viability and (2) the ability of surviving meristems to develop shoots and plantlets. Lack of regrowth often contributes to cryostorage failure and this has implications for

genebank efficiency. The arrested growth of meristems is associated with inherent adaptations induced by cryogenic and *in vitro* treatments. For potato, cryopreservation may stimulate genotype dependent dormancy which impairs meristem development during recovery. This is observed in various growth regulator and additive (BSA) treatments applied to cvs 'Golden Wonder' and 'Desiree'. Developmental time courses (Fig. 5, Table 1) were strongly influenced by recovery medium with differences of ca. 100 days in the mean time taken for plantlets to regenerate and mature (Harding and Benson 1994, Harding 1997). The dynamics of recovery and significance of physiological uniformity is manifest as synchronicity between individual potato plants regenerated within the same batch of cryopreserved meristems (Fig. 5). Recovery on TBZ medium produced the most survivors, developed the most shoots and matured the most plants, within the shortest time. In contrast, recovery on hormone free (HF) medium supported a low number of survivors, shoots and plantlets demonstrating post-cryopreservation development can be greatly improved by growth regulators. Pre- and post- cryopreservation light treatments were also found to influence survival and shoot development in recovering cryopreserved potato meristems (Fig. 1). Thus, non-cryogenic factors can be optimized to enhance survival and development after cryopreservation. Other hitherto unknown factors may also contribute to differences between shoot meristems and plant culture cycles accounting in part for inter-experimental and intra-replicate variability (Fig. 3). Similar observations are reported for the initial survival, regrowth and plantlet development of garlic following cryopreservation (Keller 2005). Differences in physiological status of meristems may also explain variation between experiments and replicate samples (Fig. 4) in diverse potato species (Golmirzaie and Panta 2000) and old varieties of potato (Schäfer-Menuhr et al. 1997). The basis of these outcomes in clonally propagated crops such as potato remains speculative but cumulative evidence suggests it is partly due to meristem physiology. Sensitivity of shoot tips from 1–2 month old *in vitro* potato plants to droplet-vitrification was related to physiological state. A progressive decline in recovery occurred

with decreasing bud ranking order from the apex (Halmagyi et al. 2005). The significance of shoot location and subculture duration of the mother plant has been similarly noted by Yoon et al. (2006) who also used droplet vitrification. Whereas, little or no differences were observed with respect to nodal segments or days of culture for potato shoots cryopreserved using encapsulation/vitrification (Hirai and Sakai 1999). These observations illustrate that intrinsic variability *in vitro* cultures does not ensure a sufficient level of physiological uniformity in some donor germplasm.

In Vitro Ageing

Sultan (2000) proposes a single genotype can produce different phenotypes in response to different *in vivo* environments and Munné-Bosch (2007) suggests intrinsic changes during the ageing of perennials become fixed in development and this affects the physiology of the whole plant. These observations are relevant to *in vitro* tissue cultures maintained for long periods, as *in vitro* ageing may differentially affect their response to cryopreservation. Age strongly influences the survival and regeneration of cryopreserved potato meristems; this is independent of genotype as older cultures are less able to recover from cryostorage (Fig. 2). Long-term *in vitro* cultures of cv 'Desiree' introduced in 1980 were unable to regrow following cryopreservation (Keller and Dreiling 2003) and 'old' and 'young' *in vitro* garlic cultures respond differentially (Keller 2005). Prolonged culture *in vitro* is known to cause loss of totipotency, vigour and promote neoplastic progression (Benson 2000, Gaspar et al. 2002) illustrating that changes in explant quality as a result of *in vitro* ageing may exacerbate genotypic responses to cryopreservation. It is therefore, recommended younger, newly initiated cultures are preferentially selected for cryopreservation or older cultures are regenerated/rejuvenated before their meristems are cryobanked.

An Oxidative Stress Model for Different Survival Responses

Recovery data of potato and *Ribes* (Figs. 1, 2, 6, 7) initially show high levels of survival in the more tolerant genotypes followed by a progressive decline which is more pronounced in sensitive genotypes. The recurrent loss of original survivors coupled with failure of those that are remaining to develop shoots results in substantial diminishing returns from the cryobank. Recovery and oxidative stress profiles of cryopreservation tolerant *R. nigrum* and sensitive *R. ciliatum* are markedly different and comprise two distinct phases (Johnston et al. 2007). Initially (1–2 weeks) up to 70% of *R. ciliatum* shoot meristems survived but by 3–5 weeks most had perished and only 20% of those remaining developed shoots (Fig. 6). In contrast *R. nigrum* meristems maintained relatively high levels of recovery throughout cryopreservation, up to 70% of the meristems survived liquid nitrogen of which ca. 100% produced shoots (Fig. 6). Oxidative stress profiles (Table 3) also revealed different genotypic responses to acclimation and cryopreservation leading to the postulation that two different oxidative stress pathways occur in *Ribes* shoots recovering from cryopreservation (Fig. 7). The first involves a deleterious reaction to oxidative stress manifest as free radical mediated (photo) oxidation, failure of antioxidant protection, degeneration, death of early survivors and loss of morphogenetic competence. Alternatively, the second pathway comprises a beneficial, programmed oxidative burst, punctuated by enhanced $\bullet\text{OH}$ production and the putative involvement of Reactive Oxygen Species (ROS) in anti-stress signaling. This results in an elevated antioxidant status, enhanced tolerance to cryopreservation and the conversion of ca. 65–80% of the cryopreserved meristems to shoots (Fig. 7). These differential responses were initiated during sucrose-simulated cold acclimation and subsequent tolerance to liquid nitrogen correlated to increases in $\bullet\text{OH}$ activity, C_2H_4 evolution, antioxidant status, phenolic accumulation, protein SH group status and a reduction in chlorophyll and carotenoid photooxidation as compared to the sensitive genotype (Table 3, Fig. 7). Importantly,

antioxidant protection in *R. nigrum* persisted during recovery indicating elevated antioxidant status is associated with cryopreservation tolerance (Johnston et al. 2007). These findings concur with *in vivo* studies of freezing stress in which antioxidant protection acquired during cold acclimation and drought tolerance is associated with genotypic tolerance to freezing (Blödner et al. 2005, Dale and Heiberg 1984, Mather et al. 1980).

Is Programmed Cell Death a Factor in Delayed Post-Cryopreservation Failure?

Baust (2002) proposes delayed onset, cryopreservation-induced cell death as a significant contributory factor in storage failure and recommends that apoptosis and necrosis should be considered when developing biopreservation protocols. As a delayed decline in recovery has been observed in plant meristems following cryopreservation (Figs. 1, 6) there may be some parallels with animal tissue cryostorage. Cell death incurred by cryoinjury is known to be associated with a belated free radical cascade in human tissues (Hoffman and Bischoff 2004) and apoptosis is induced *via* the release of cytochrome *c* from mitochondria (Reape and McCabe 2008). Thus, oxidative stress and apoptosis may become causal and contributory factors in predisposing both plant and animal tissues to immediate and delayed onset cryoinjury. In evaluating programmed cell death in plant cryopreservation it is important to differentiate between events that lead to necrosis and those that are confirmed as truly apoptotic by using biochemical markers (Bischof et al. 2002, Gage and Baust 2002, Reape et al. 2008). As apoptosis remains to be fully elucidated in plants, it is also necessary to distinguish between necrosis and apoptosis-like programmed cell death during cryostorage recovery responses (Reape and McCabe 2008). Nevertheless, understanding the mechanisms involved is important as degenerative responses are often observed in storage sensitive genotypes and the delayed loss of survivors can be sufficiently significant to affect the efficiency of plant genebank operations. The fact that the delayed

decline in meristem survival is frequently independent of protocol suggests cryopreservation stresses *per se* may exacerbate degenerative pathways, some of which proceed *via* necrosis, whilst others may involve programmed cell death (Reape et al. 2008). In the case of certain genotypes, optimisation of cryogenic factors may not be sufficient and other strategies will be needed to ensure sustained recovery. These are likely to include the manipulation of physiological status prior to cryopreservation, acclimation treatments, choosing vigorous, young donor cultures and excising meristems in an appropriate stage of development. Exploring apoptotic induction factors in both immediate and delayed onset post-storage degeneration in plant germplasm has practical merits particularly as it is under investigation in other biorepository sectors (Khan et al. 2009, Baust 2002).

Cryoinjury, Genetic Stability and Cryobionomics

Physiological factors are likely to contribute to variable responses independent of the cryopreservation protocol and an investigative molecular approach may help elucidate the basis for these differences. Figure 5D and Table 2 show rDNA fragment variability within a given batch of individual cryopreserved plants providing further support for the premise that physiological variation occurs during *in vitro* manipulation (Harding et al. 2008, Carpentier et al. 2007). Ribosomal fragment variability possibly results from restriction enzyme sensitivity to methylation of enzyme recognition sites (Harding 1997). DNA methylation as an epigenetic mechanism for genomic variability (Harding 2004, Harding et al. 2008) may also account for physiological variation between individual meristems and their differential responses (Johnston et al. 2007, Johnston et al. 2009). Although, *in vitro* variation within batches of individual meristems clearly exists, it is not inherently obvious prior to cryopreservation and this makes its detection difficult. The use of DNA methylation as an epigenetic marker for physiological variability may be useful in evaluat-

ing recalcitrant germplasm before cryopreservation (Johnston et al. 2005, Harding et al. 2008, Johnston et al. 2009). There is evidence that differences between plants showing differential methylation affect their regeneration and maturation following cryopreservation (Fig. 5, Harding et al. 2008). The time taken for *in vitro* plantlets (E, F) to regenerate is also related to differential rDNA methylation as evidenced by the delays incurred on hormone free recovery medium compared to plantlets (I, J) with similar levels of methylation and recovery times (Fig. 5A) using more complex medium (Harding 1997). Although, the respective positions for the plantlets (I, J) are relatively unchanged (Fig. 5A, B) others (E, F) show the opposite trend in recovery times between growth *in vitro* and maturation (Harding and Benson 1994). Clearly, there are complex interactions between the epigenetic regulation of gene expression and hormone induced physiological responses in these plants.

Collective evidence illustrates the intricate combined effects of cryogenic and non-cryogenic factors following the exposure of shoot meristems to the physical, chemical and physiological stresses that cause cryoinjury (Johnston et al. 2007, Johnston et al. 2009). The effects of cryoinjury upon the genome are often unknown; any accumulative DNA polymorphisms may not be induced by solely cryogenic factors but result from the whole process (Harding 2004). The intrinsic cellular and metabolic factors (Carpentier et al. 2007, Harding et al. 2008) that affect physiological status can profoundly determine meristem survival. For example, where, survival is mostly dependent on antioxidant defences that oppose the deleterious effects of cryoinjury (Johnston et al. 2007). The hypothetical relationship between cryoinjury, genetic stability and cryobionomics is shown in Figure 8 which highlights the potential impacts of cryoinjury on the genome, transcriptome, proteome and metabolome. This leads to several possible changes, those incurred by modulated patterns of gene expression mediated through epigenetic mechanisms because of naturally adaptive responses, as well as events that lead to undesirable instability. These may be manifest as detectable changes that affect trueness-to-type with respect to morphology, histology, cy-

Harding, K. Exploring cryopreservation in clonally propagated crops

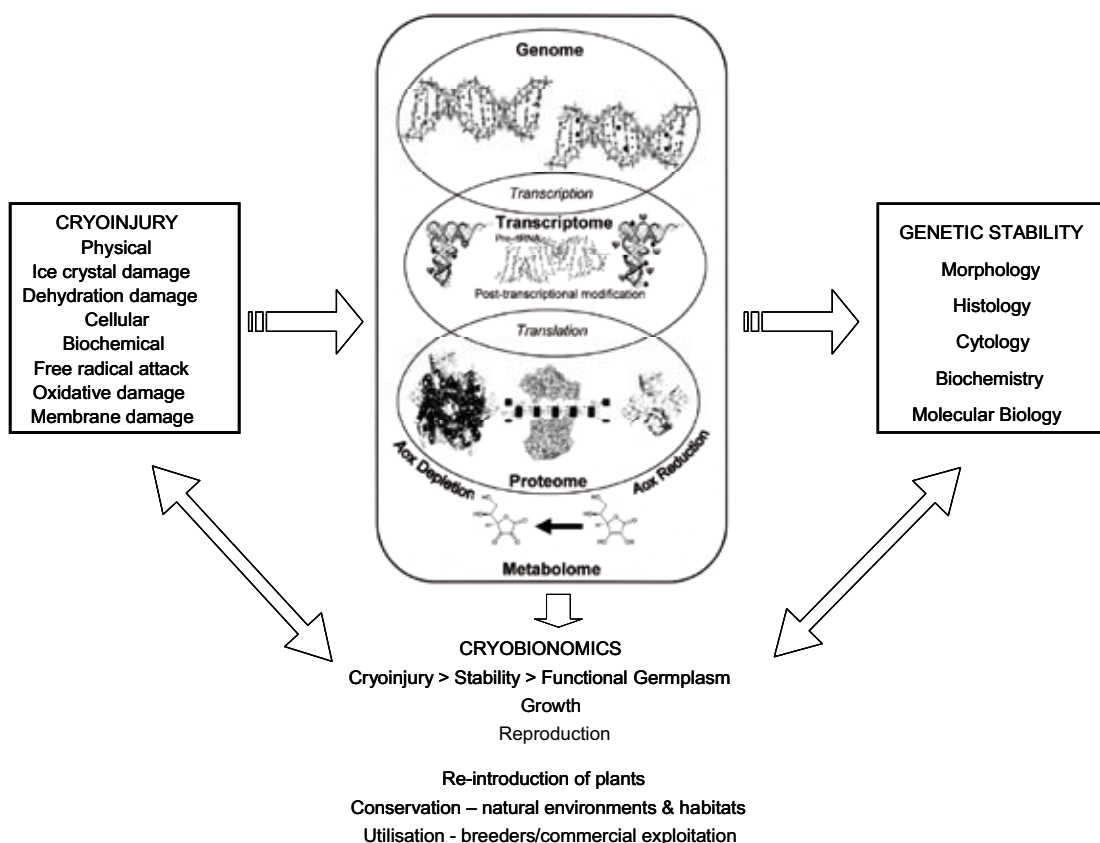


Fig. 8. Relationship between cryoinjury, genetic stability and cryobionomics.

tology, biochemistry and the molecular biology of whole plants recovered from cryopreserved germplasm (Harding 2004). Profound epi- and/or genetic alterations to the genome considered within the framework of ‘cryobionomics’ are likely to affect the function of germplasm possibly by disrupting established patterns of growth and reproduction. These changes may influence both the quality of conserved germplasm and the re-introduction of plants recovered from cryobanked materials into natural environments and habitats, and including their utilisation by breeders and exploitation in commercial activities. This study recommends a more

holistic approach is now required to understand the basis of success or failure following cryostorage. Cryobionomics provides a conceptual framework to explore the linkages between cryogenic and non-cryogenic stress factors by using molecular-physiological approaches to aid cryopreservation protocol improvement and optimization.

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