

Asymmetric Somatic Plant Hybridization: Status and Applications

Lakshmanan Prabhu Shankar^{1,2}, Eeckhaut Tom¹, Deryckere Dieter¹, Van Bockstaele Erik^{1,2}, Van Huylenbroeck Johan¹

¹Institute for Agricultural and Fisheries Research (ILVO), Plant Sciences Unit, Applied Genetics and Breeding, Melle, Belgium;

²Department of Plant Production, Faculty of Bioscience Engineering, Ghent University, Ghent, Belgium.

Email: johan.vanhuylenbroeck@ilvo.vlaanderen.be

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ABSTRACT

To create asymmetric somatic hybrids, the genome of the so-called donor protoplast is fragmented prior to protoplast fusion. As a result, only a limited amount of the donor genome is transferred to the fusion product. This technique can circumvent some commonly observed problems related to symmetric fusion and offers a practical breeding tool for asexual hybridization. Genomes are typically fragmented by irradiation, microprotoplast production or application of metabolic inhibitors such as iodoacetamide. Irradiation and microprotoplast production fragment the nuclear genome, whereas iodoacetamide inactivates the cytoplasmic genome. It can therefore be used to introduce cytoplasmic male sterility, an important practical application. For hybrid verification and genome characterization, molecular markers and cytogenetic techniques are applied. This review highlights and discusses progress made during the last decade in spermatophytes asymmetric protoplast fusion.

Keywords: Asymmetric Somatic Fusion; Genome Fragmentation; Protoplast

1. Introduction

Plant cells from which the cell wall has been enzymatically or mechanically removed are called protoplasts. Theoretically, protoplasts are totipotent, meaning that after their isolation and subsequent culture they have the capability to dedifferentiate, re-start the cell cycle, go through repeated mitotic divisions and then proliferate or regenerate into various organs. Fusion of protoplasts from different species can therefore be a practical breeding tool [1] and circumvents sexual hybridization related prezygotic or postzygotic barriers [2]. Somatic hybridization differs from other techniques in many respects. When comparing somatic hybridization to transgenic approaches, somatic hybridization enables broadening of the germplasm base, allows the transfer of uncloned multiple genes and generates products that are not subjected to the same legal regulations as transgenic lines [3,4]. Also, it transfers both mono- and polygenic traits [5]. The first protoplast fusion was described more than a century ago by Küster [6]. Tobacco was the first crop in which successful interspecific somatic hybridization was

reported [7]. Since then, improvements and somatic hybrids have been made in many species and complete plant regeneration was accomplished. A detailed historical overview of breakthroughs in protoplast related research is incorporated in Gamborg [8] and Davey *et al.* [9].

Protoplast fusion can be either symmetric or asymmetric depending on the nature of genetic contribution of the fusion partners. A somatic breeding protocol can typically be subdivided into the following steps: isolation, fragmentation (in case of asymmetric hybridization), fusion, regeneration and selection [10,11] (**Figure 1**). In symmetric fusion the complete genomes of both parent protoplasts are fused. However, when two complete genomes fuse, a phenomenon called “gene conflict” may arise, because certain chromosomes repel one another. Moreover, the technique introduces a significant amount of unwanted genetic material. These limitations result in abnormal growth, regeneration of hybrids with low fertility, non-rooted shoots, slow hybrid growth, and recalcitrant calli or microcalli [2].

In asymmetric fusion, after genome fragmentation only a limited amount of one genome is transferred to the

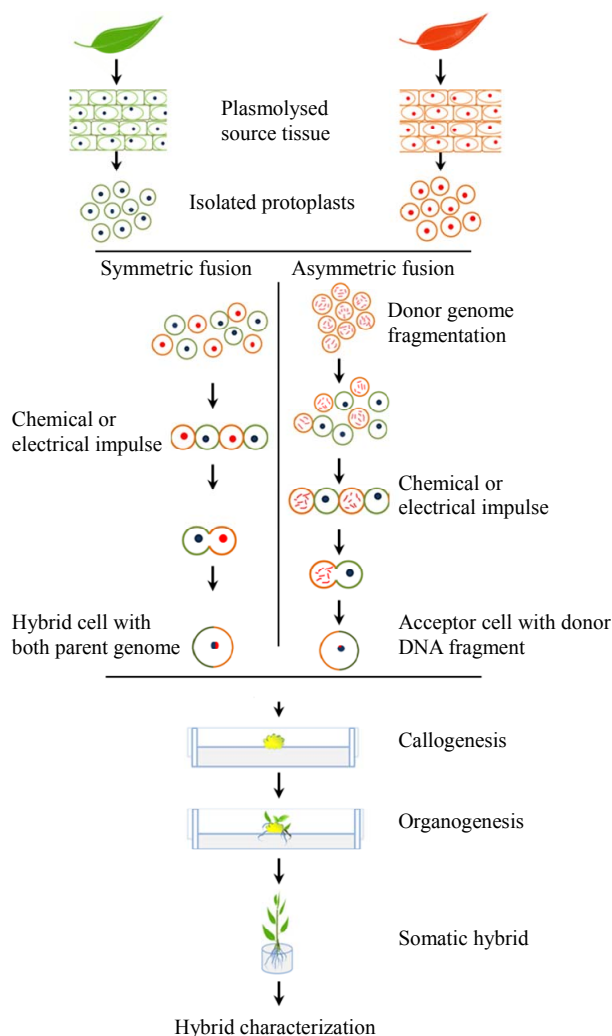


Figure 1. Schematic representation of symmetric and asymmetric somatic hybridization techniques.

fusion product [12,13] (**Figure 1**). Cytoplasmic genomes can also be recombined with nuclear genomes for applications such as cytoplasmic male sterility (CMS) introduction [11]. The technique is being applied to circumvent the above mentioned barriers in symmetric fusion. For example, symmetric hybrids between *Brassica napus* and *Lesquerella fendleri* are self-sterile, whereas asymmetric hybrids between the same fusion parents are self-fertile [14]. Similarly, symmetric fusion between *Orychophragmus violaceus* and *B. napus* yields sterile hybrids, whereas asymmetric hybrids are fertile and can set seeds [15]. By introgressing fewer genes than after sexual crossing or symmetric somatic fusion, the number of backcrosses can also be significantly reduced. Genome fragmentation of the donor parent encourages the elimination of much of its redundant genetic material in the somatic hybrid. Moreover, in asymmetric fusions, most karyotype instability causing donor genes are eliminated during the first post-fusion mitoses, as opposed to sym-

metrical fusions after which eliminations can occur up to the first sexually derived generation [16]. In other words, not only does asymmetric fusion introduce fewer genes in a recipient genome after fragmenting the donor genome, but elimination of disadvantageous genes or chromosomes also proceeds quicker. Nevertheless, chromosomes and chromosome fragments can still be lost during meiosis due to rearrangements [17].

The objective of this review is to provide an overview of the development of technologies for fragmentation and screening. Besides the progress made in spermatophytes asymmetric protoplast fusion related research during the last decade is highlighted and discussed.

2. Fragmentation Techniques

In order to obtain asymmetric fusions, only partial genomes are transferred. Several techniques can be used for genome fragmentation of the so-called “donor” genome: irradiation, microprotoplasts (MPPs), or metabolic inhibitors as iodoacetamide (IOA). In some cases asymmetric fusions were realized without fragmentation treatment. The production of these asymmetric hybrids decreased possible long term irradiation effects on hybrid growth and development [18].

2.1. Irradiation

Genome fragmentation by irradiating can be achieved by using either ionizing radiation techniques (X or gamma rays) or non ionizing radiation (UV rays). Irradiation often induces random chromosome breakage, but also gene deletion and rearrangement and can be responsible for hybrid sterility [19]. The first application of X-rays to obtain asymmetric hybrids was performed in parsley [20]. UV light used to create asymmetric hybrids was used for the first time on *Nicotiana* donor protoplasts [21]. Earlier, X or gamma rays were more frequently used for donor protoplast fragmentation, but nowadays UV treated protoplasts are more widely applied. Both irradiation types efficiently induce asymmetric somatic hybrids in a dose-dependent manner, e.g., between *Brassica napus* and *Arabidopsis thaliana* [22]. Nonetheless, within a single species susceptibility towards different irradiation types can vary strongly [23]. Hall *et al.* [24] investigated whether UV radiation could be used as an alternative for ionizing radiation techniques. They found that UV had a detrimental effect on sugar beet protoplasts: resynthesis of a cell wall, cell growth and cell division were partially or totally eliminated. However, protoplast viability had not decreased after 6 days culture, but after 14 days, the UV treated cells died. On the other hand, a significant advantage of UV radiation over ionizing radiation was its easy application and high reproducibility. Similar observations were made after exposure of *Cichorium* proto-

plasts to UV [25]. In cucumber, the negative effect of UV-C irradiation on cell wall regeneration, protoplast viability and the intensity of the nuclei after DAPI staining was also demonstrated [26].

A general problem is the quantification of DNA damage after an irradiation treatment. Abas *et al.* presented Comet assay single cell gel electrophoresis (SCGE) as a reliable tool to observe single and double strand breaks in mesophyll protoplasts of *Nicotiana plumbaginifolia* [27]. Xu *et al.* revealed extensive DNA fragmentation in UV irradiated *Citrus unshiu* protoplasts with the terminal deoxynucleotidyl transferase biotin-dUTP nick end labeling assay [28]. However, both methods are not generally applicable.

2.2. Microprotoplast Mediated Chromosome Transfer

Next to donor protoplast irradiation, micronuclei and microprotoplasts mediated chromosome transfer (MMCT), which was originally developed for mammalian cells, has been considered as an alternative method for partial genome transfer [29].

Mass induction of micronucleation and efficient isolation of microcells are key steps in any MMCT for successful transfer of partial genomes [30]. Microtubules are involved in several processes such as chromosome migration, cell structure, cellulose microfibrils guidance and arrangement, cell wall formation, intracellular movement and cell differentiation. Toxic substances as anti-mitotic herbicides or colchicine prevent their normal polymerization [31]. Application of these spindle toxins to synchronized cells generally blocks cells in metaphase and scatters chromosomes in the cytoplasm; subsequently, those decondense into micronuclei. Subsequently, these micronucleated cells are stripped of their cell wall, and the resulting MPPs are ultracentrifuged to subdivide them into classes [32]. These can be further filtered through sequential filters of smaller pore width. Some recent examples of suspension cell derived MPPs are *Citrus unshiu* [33] and *Beta vulgaris* [34].

In developing microspores of ornamental species like *Lilium* and *Spathiphyllum*, micronuclei were induced through the action of mitosis arresting chemicals, without synchronization requirement [35,36]. By using microspores instead of suspension cultures the risk of mutation accumulation in suspension cells can be avoided [37].

Regardless of the source material, the efficiency of spindle toxins depends on their type, dose, incubation period and the plant genotype. Those parameters can be optimized, as recently demonstrated for *Spathiphyllum wallisii* [36]. For this crop, the highest micronuclei indices were obtained after microspore treatment with 10 μ M oryzalin for 72 h or 20 μ M chlorpropham for 48 h for a particular model genotype. The maximal number of mi-

cronuclei observed was 12, while the haploid chromosome number amounts 15. Oryzalin is the most widely used mitosis inhibitor, but its efficiency varies based on the plant species [38,39]. Famelaer *et al.* quantified genome fragmentation in *Beta* microprotoplasts through flow cytometry and confocal microscopy [34].

2.3. Cytoplasmic Inactivation

Metabolic inhibitors, such as IOA and rhodamine 6-G can be used to obtain asymmetric fusions. The exact mode of action of IOA has not yet properly been described, the compound however inhibits protoplast division by irreversibly inactivating the cytoplasm. In red chicory mesophyll protoplasts division was totally inhibited after treatment with 2 - 4 mM iodoacetate [40]. When using IOA, a lower optimal concentration of 1.625 mM was found [25]. The lower optimal IOA concentrations compared to iodoacetate might be explained by better cell penetration of IOA. Similar optimal IOA concentrations were found in *Musa* [41]. Lower IOA concentrations (0.5 mM) stopped growth of *Gossypium hirsutum* protoplasts, whereas 3 mM and 7.5 mM IOA were required to stop cell proliferation in *Citrus* [42] and *Bupleurum scorzoniferifolium* [43], respectively.

Fusion of IOA-treated recipient parental protoplasts with irradiated donor protoplasts could produce cybrids. In *Cichorium*, successful asymmetric protoplast fusion has been performed between γ -rays-irradiated sunflower protoplasts and iodoacetate-treated red chicory protoplasts [40]. IOA treatments prevent cell division, but fusion with non-IOA treated protoplasts restores cell division ability, thus opening ways towards heterokaryon or cybrid selection.

3. Asymmetric Hybridization

In earlier protoplast reviews, the concept of asymmetric hybridization is well explained, but has not been the focus of the review [11,12]. Fragmentation techniques have been highlighted, however, [24,29] also in practical manuals [44]. In the review by Xia [13], wheat is used as an example to discuss chromosome engineering through asymmetric fusion.

Over the last decade, asymmetric fusion techniques have been widely applied and several new asymmetric hybrids were obtained (**Table 1**). The most studied families were Brassicaceae and Poaceae, followed by Rutaceae. The number of asymmetric hybridization realized through PEG fusion was 4-fold the number of fusions generated by electrical fusion.

Asymmetric hybridization has allowed new genome combinations that would be more difficult, if not impossible, to realize through classical symmetric fusion or sexual crossing. For the first time, an asymmetric hybrid

Table 1. Progress in asymmetric protoplast fusion in different plant families during last decade (2004-2013).

Plant family and species (acceptor + donor) [†]	Aim	Cell source ^u	Fragmentation tool ^v	Fusion tool ^w	Characterization			Ref.
					Cytogenetic ^x	DNA markers	Other methods	
Apiaceae + Gentianaceae								
<i>Bupleurum scorzonerifolium</i> + <i>Swertia mussottii</i> *	Secondary metabolites	C	UV	PEG	GISH	RAPD, SQ RT-PCR	Isozyme analysis, mitochondrial and cpDNA specific probes on Southern blots, HPLC	[45]
<i>Bupleurum scorzonerifolium</i> + <i>Swertia tetraptera</i> *	Secondary metabolites	SC + C	UV	PEG	CC, GISH	RAPD, SSR(C)	Isozyme analysis, HPLC	[46]
<i>B. scorzonerifolium</i> + <i>Gentianopsis paludosa</i> *	Secondary metabolites	SC	UV	PEG	CC	RAPD, RFLP	Isozyme analysis, HPLC, GC-MS, 5S rDNA spacer sequence analysis	[47]
Asteraceae								
<i>Helianthus annuus</i> + <i>H. maximiliani</i> *	Biotic resistance	H + M	UV	EF		RAPD	Isozyme analysis	[48]
Brassicaceae								
<i>Brassica napus</i> + <i>Isatis indigotica</i> *	Genetic variation, Secondary metabolites	M	IOA(A) + UV	PEG	CC, GISH	AFLP, CAPS(C) ^y	Pollen fertility	[49]
<i>Brassica napus</i> + <i>Orychophragmus violaceus</i>	Chromosome addition lines	M	IOA(A) + UV	-	CC, GISH			[50]
<i>Brassica oleracea</i> + <i>B. nigra</i>	Genetic variation, Biotic resistance	H + M	UV	PEG	CC, FCM, GISH	AFLP, CAPS(C), CAPS(M) ^z	MtDNA specific probes on Southern blots, resistance screening	[51]
<i>Brassica oleracea botrytis</i> + <i>B. carinata</i> + <i>B. juncea</i> + <i>B. nigra</i> + <i>Sinapis alba</i>	Biotic resistance	H + M	X	PEG		RAPD	Resistance screening	[52]
<i>Brassica oleracea capitata</i> + <i>Barbarea vulgaris</i> + <i>Capsella bursapastoris</i> + <i>Diplotaxis tenuifolia</i> + <i>Hesperis matronalis</i> + <i>Matthiola incana</i> + <i>Raphanus sativus</i> + <i>Sinapis alba</i>	Biotic resistance	H + M	X	PEG		RAPD	Resistance screening	[52]
<i>Orychophragmus violaceus</i> + <i>Lesquerella fendleri</i> (GFP)*	Plastome transfer	M + C	γ	PEG		ITS, CAPS(M)	Isozyme analysis, GFP	[53]
Brassicaceae + Apiaceae								
<i>Arabidopsis thaliana</i> + <i>Bupleurum scorzonerifolium</i> *	Secondary metabolites	C	IOA(A) + UV	PEG	CC, GISH,	RAPD	5S rDNA spacer sequence analysis	[43]
<i>Arabidopsis thaliana</i> + <i>Bupleurum scorzonerifolium</i>	Fragmentation tool	-	UV	PEG	CC, GISH	RAPD, SSR		[54]
<i>Arabidopsis thaliana</i> + <i>Bupleurum scorzonerifolium</i>	Hybrid analysis, fragmentation tool	C + SC	γ	PEG	CC	RAPD, SSR	Histology	[23]
Brassicaceae + Poaceae								
<i>Arabidopsis thaliana</i> + <i>Triticum aestivum</i> *	Hybrid analysis	SC + C	UV	PEG	CC, GISH	RAPD, SSR, CAPS(C)	Isozyme analysis	[55]
Malvaceae								
<i>Gossypium hirsutum</i> + <i>G. klotzschianum</i>	Alternative for symmetric somatic hybridisation	SC	UV	EF	CC	RAPD, SSR, CAPS(C)		[56]
Musaceae								
<i>Musa</i> "Guoshanxiang" + <i>M. acuminata</i> *	Biotic resistance	SC	IOA(A) + UV	PEG	CC	RAPD, ISSR		[41]

Continued

Poaceae								
<i>Festuca arundinacea</i> + <i>Triticum aestivum</i> *	Hybrid analysis	SC	UV	PEG	CC, GISH	RAPD, SSR(C), MSAP	Isozyme analysis, mtDNA specific probes on Southern blots	[57]
<i>Oryza sativa japonica</i> + <i>O. meyeriana</i> *	Biotic resistance	SC	IOA(A) + X	PEG	CC	RAPD	Resistance screening	[58]
<i>Triticum aestivum</i> + <i>Avena sativa</i>	Hybrid analysis	SC	UV	-	ISH	SSR		[59]
<i>Triticum aestivum</i> + <i>Haynaldia villosa</i>	Biotic resistance, protein content	SC + C	γ	PEG	CC, GISH	RFLP(C)	Isozyme analysis, 5S rDNA spacer sequence analysis	[60]
<i>Triticum aestivum</i> + <i>Lolium multiflorum</i> *	Biotic resistance	SC	UV	-	CC	RAPD, SSR	Isozyme analysis, mtDNA specific probes on Southern blots	[61]
<i>Triticum aestivum</i> + <i>Lolium multiflorum</i>	Agronomic traits, biotic resistance	SC	IOA(A) + X	EF	CC	RFLP, AFLP	Isozyme analysis, mtDNA specific probes on Southern blots	[62]
<i>Triticum aestivum</i> + <i>Lolium multiflorum</i>	Radiation hybrid panel/ genome mapping	-	UV	-	GISH	RFLP, SSR	Sequencing	[63]
<i>Triticum aestivum</i> + <i>Setaria italica</i> *	Abiotic resistance	(SC + C) + C	UV	PEG	GISH, CC	RAPD, RFLP(C), RFLP(M)	Isozyme analysis, 5S rDNA spacer sequence analysis	[64]
Poaceae + Apiaceae								
<i>Festuca arundinacea</i> + <i>Bupleurum scorzonerifolium</i> *	Hybrid analysis	SC	UV	PEG	CC	RAPD	Isozyme analysis, 5S rDNA spacer sequence analysis	[65]
<i>Triticum aestivum</i> + <i>Bupleurum scorzonerifolium</i> *	Genetic variation, genome mapping	C + SC	UV	PEG	GISH	CAPS, RAPD, SSR	Isozyme analysis	[66]
Rutaceae								
<i>Citrus paradisi</i> + <i>C. sinensis</i>	Genetic variation	SC	IOA(A) + γ	-	FCM	AFLP		[42]
(<i>Citrus reticulata</i> x <i>C. sinensis</i>) + <i>C. sinensis</i>	Genetic variation	SC	IOA(A) + γ	-	FCM,	AFLP		[42]
<i>Citrus sinensis</i> + <i>C. unshiu</i>	Fragmentation tool	SC	UV	EF	CC, FCM	RAPD, AFLP, CAPS(C)		[28]
Solanaceae								
<i>Nicotiana tabacum</i> + <i>N. repanda</i>	CMS	M	Rhodamin 6G (A)	PEG	CC	RAPD, CAPS(M)	Isozyme analysis	[67]
<i>Petunia hybrid</i> + <i>Nicotiana tabacum</i>	Plastome transfer	M	UV	PEG		RAPD, CAPS(C)	MtDNA specific probes on Southern blots, chloroplast RNA specific probes on Northern blot	[68]

*Species labeled with * were fused for the first time; ^uC: callus; CO: cotyledon; H: in vitro hypocotyls; M: in vitro mesophyll cells; SC: suspension cells; - : described in earlier publications; ^v(A): for acceptor; IOA: iodoacetamide; UC: ultracentrifugation; UV: UV ray irradiation; X: X ray irradiation; γ : gamma-ray irradiation; ^wEF: electrofusion; PEG: chemical fusion with polyethylene glycol; - : described in earlier publications; ^xCC: chromosome counting; FCM: flow cytometry; (G) ISH: (genomic) in situ hybridization; ^y(C) on cpDNA; ^z(M) on mtDNA.

was reported in banana [41]. Interfamilial asymmetric hybrids have been produced between the dicot *Arabidopsis thaliana* and the monocot common wheat [55]. Fusion between phylogenetically remote tall fescue, Italian ryegrass and common wheat was achieved [57,61,62]. In cotton, asymmetric hybrids were obtained as an alternative for symmetric hybrids [56]. New somatic hybrids were obtained between monocot *Festuca arundinacea* and dicot *Bupleurum scorzonerifolium* through symmetric as well as asymmetric fusions [65]. The first successful somatic hybrid regeneration between *Oryza sativa* L. ssp. *japonica* and *O. meyeriana* L. was reported [58].

Scholze *et al.* produced the first *Raphanus-Brassica* somatic hybrids with fungal and virus disease resistance [52]. Cybrids were produced between chloroplast transformant tobacco and petunia [68]. Using UV irradiated asymmetric hybrids a radiation hybrid panel was established for *Lolium multiflorum* [63]. Taski-Adjukovic *et al.* [48] regenerated an asymmetric hybrid between sunflower and *Helianthus maximiliani* for the first time.

Acceptor protoplast sources for asymmetric hybridization existed mainly of suspension cell cultures, mesophyll, callus and hypocotyls. The donor protoplast source differed in about 30% of the cases from the one for the

acceptor. Brassicaceae and Asteraceae hypocotyl acceptor protoplasts were combined with mesophyll donor protoplasts [48,51,52].

Biotic resistance introduction, genetic variation, agronomic traits such as seedless fruits, hybrid analysis, fragmentation technology development and secondary metabolite production were the most important recent aims for asymmetric hybridization. Abiotic resistance introduction, hybridization, genome mapping and the establishment of chromosome addition lines were rare objectives (**Table 1**). In *Bupleurum scorzonerifolium*, asymmetric hybrids were obtained after protoplast fusion of UV treated *B. scorzonerifolium* and wheat protoplasts. However, instead of generating wheat carrying *B. scorzonerifolium* chromosome fragments, the reverse was found. This study can be of major interest for the construction of physical maps of the wheat genome [66]. The same was observed when untreated *Arabidopsis thaliana* protoplasts were fused with UV treated *Bupleurum* protoplasts [43].

Other motives were plastome and/or CMS transfer [67]. The latter is an important practical application of new genome/cytoplasmic combinations. Fitter *et al.* [69] demonstrated the possibility of introgressing CMS carried by mtDNA from a wild species into the cultivated crop. In *Cichorium*, CMS was introduced after asymmetric fusion with sunflower [40]. Sheahan *et al.* [70] reported the phenomenon of massive mitochondrial fusion (MMF) which leads to near-complete mixing of the mitochondrial population within 24 h. MMF appears specific to dedifferentiation, since it also occurs in mesophyll protoplasts of *Arabidopsis* and *Medicago* but not in protoplasts from already dedifferentiated cells such as tobacco BY-2 or callus cultures. These results allow a clearer interpretation of how novel mitochondrial genotypes develop following cell fusion. In other investigations, Sytnik *et al.* [71] demonstrated that also chloroplasts can be transferred to remote species by protoplast fusion.

4. Characterization of Asymmetric Somatic Hybrids and Genome Stability

Apart from morphological characterization, the majority of the publications reporting on complete plant regeneration describe the use of molecular tools to unravel the genomic constitution of the alleged hybrids (**Table 1**). DNA markers were sometimes complemented with isozyme analysis, sodium dodecyl sulphate polyacrylamide gel electrophoresis or sequence analysis. The most frequently employed molecular markers were Random Amplification of Polymorphic DNA (RAPD), Simple Sequence Repeats (SSR), Amplification Fragment Length Polymorphism (AFLP), Restriction Fragment Length Polymorphism (RFLP) and Cleaved Amplified Poly-

morphic Sequence (CAPS). PCR-RFLP and CAPS analysis using mitochondrial or chloroplast universal primer pairs were efficient and reliable methods for characterizing the cytoplasmic genome. This technique was applied for both chloroplast and mitochondria screening, whereas SSR was only used once for chloroplast evaluation. Southern blotting for cpDNA and mtDNA was employed to screen cytoplasmic DNA, whereas Northern blotting was used once for chloroplast evaluation. Compared to RFLP with labeled probes, CAPS is simpler, more rapid and less expensive [72]. Chloroplast SSR is even more convenient and efficient since enzyme cutting following PCR reaction is not needed [73]. Also sequencing of common bands and searching for restriction endonuclease sites could be cheaper and more convenient than actual CAPS analysis (though after sequencing CAPS could be used to confirm the results).

Besides molecular markers, cytogenetic tools as chromosome counting, flow cytometry and genomic in situ hybridization (GISH) can be used to distinguish asymmetric hybrids (**Table 1**). Especially the latter technique enables to visualize the hybrid genomic constitution and to follow genomic stabilization. After a symmetric fusion of two *Triticum* genotypes with *Psathyrostachys*, GISH analysis showed that the hybrids were highly asymmetric and contained only wild rye chromosome fragments, whereas the strong relationship of the hybrids and wheat was demonstrated by SSR markers [18]. Also the chloroplasts of the hybrids and wheat were identical. Apparently the genetic complementation of 2 *Triticum* parents stimulates the rejection of wild rye donor chromosomes. In the asymmetric *Triticum aestivum* + *Setaria italica* fusion, genome complementation can be used as a selection tool. The only regenerative callus type has 5 recombinant chromosomes and a chromosome count of 48, exceeding the 42 of normal wheat [64]. In non-regenerative calli, only 0 - 2 *Setaria* chromosomes were present. Symmetric [74] as well as asymmetric [43] *Arabidopsis thaliana* + *Bupleurum scorzonerifolium* somatic fusions were made. The symmetric hybrids contain the complete *B.s.* chromosome set, 0 - 2 *A.t.* chromosomes and some nuclear or cytoplasmic *A.t.* fragments. Also after *B.s.* UV irradiation, surprisingly *A.t.* genes were introgressed in the *B.s.* genome whereas the opposite was expected. In other words, *A.t.* chromatin is preferentially eliminated, the type of cross merely defines whether full chromosomes or DNA fragments were integrated.

Like nuclear genomes, cytoplasmic genomes are not always stable upon fusion. Intergenomic chloroplast recombination is a rare event in higher plants in contrast to mitochondrial genomes that show high recombination levels [75]. Sequencing, used for searching restriction endonuclease sites, can be efficiently combined with CAPS to demonstrate mtDNA recombination. MtDNA

recombination was proven in *Triticum aestivum* + *Setaria italica* [64] and *Arabidopsis thaliana* + *Brassica oleracea* [76]. The latter authors proposed mitochondrial recombination as a tool for CMS introduction in cabbage. Although more rarely occurring, cpDNA recombination in hybrids has been demonstrated. In *Triticum aestivum* + *Setaria italica* hybrids, cpDNA coexistence as well as recombination occur [64]. It was also observed in *Bupleurum scorzonerifolium* + *Swertia mussottii* [46].

5. Conclusions

Recently, interest in protoplast related research has been renewed. Fusion of protoplasts allows us to move from traditional plant breeding methods to asexual methods. Compared to symmetric somatic hybridization, asymmetric fusion has some important advantages as it may limit genome conflicts. As shown in recent publications asymmetric fusion can create genomic variation for plant quality and yield improvement in agricultural or industrial crops. Especially the possibility to introduce CMS in important agricultural crops offers the breeders a tool towards hybrid seed production.

Interesting opportunities for further research may be found in the development of advanced methods for genome fragmentation including the combination of multiple techniques, such as the irradiation of MPPs, or the creation of MPPs from unreduced gametes formed by interspecific hybrids. The latter might open the possibility to transfer recombined chromosomes in a single step. Moreover, MPPs could be selected based on filtration properties, which would result in different genome types, enabling researchers to attribute plant traits to particular chromosomes and further utilize MPPs accordingly. Furthermore, the fast evolution in marker development will allow more profound studies on genome stability. Furthermore, the contribution of GISH to genome characterization studies after fusion will further increase and can be used as a tool for monitoring genomic stabilization.

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