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5	Resilience of willows (Salix spp.) differs between families during and after
6	flooding according to floodwater depth
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### 26 Abstract

27 Although the morphological and physiological responses of willows to flooding have 28 already been characterized, less is known about their responses during the post-29 flooding period. After the end of the stress episode, plants may modify some leaf and 30 plant traits to compensate for biomass loss. The aim of this work was to analyze the 31 post-flooding responses of different willow genotypes under two different depths of 32 floodwater. The hypothesis was that the growth recovery in the post-flooding period 33 would be different according to the genotype and the floodwater depth. We analyzed 34 three genotypes of five willow families (4 interspecific hybrids and one open-pollinated 35 family). The treatments were: 1) Control: plants watered to field capacity; 2) T10: water 36 covering 10 cm above soil level; 3) T65: water covering 65 cm above soil level. Both 37 flooding treatments were followed by a period of recovery (without flooding). Growth 38 was reduced by flooding in T65 but not in T10, while root-to-shoot ratio was reduced in 39 both flooding treatments. The relative growth rate in height, leaf nitrogen concentration, 40 stomatal conductance and electron transport rate changed in a different manner during 41 the post-flooding period, depending on the treatment and genetic background. These results emphasize the need for evaluating a post - flooding recovery period for the 42 43 breeding of willow genotypes destined for areas under risk of flooding. According to our 44 results, Salix matsudana could be a source of flooding tolerance for willow breeding 45 programs.

46

47 **Key words**: relative growth rate; root-to-shoot ratio; leaf nitrogen concentration

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Key message: Willows differ in their post - flooding responses according to floodwater
 depth and genotype

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## 53 Introduction

Willows (*Salix* spp.) naturally grow near riverbanks and floodplains, and they are considered as flood - tolerant forest trees (Karrenberg 2002). As a result, willow plantations can be developed in areas with high risk of flooding, either as a source of biomass, pulp and timber (Balatinecz et al. 2014), or with the purpose of restoring disturbed landscapes (Wang et al. 2017).

The morphological and physiological responses of willows to flooding have been studied extensively, and they vary according to the genotype, the length and frequency of the stress episodes, and the depth of the floodwater (Li et al. 2004, Markus - Michalczyk et al. 2016, Doffo et al. 2017, Rodríguez et al. 2018). Nevertheless, the responses of willows during the post - flooding period have received less attention (Jackson and Attwood 1996, Wang et al. 2017).

65 Global warming is expected to increase the occurrence of flooding episodes in 66 several areas of the world (Kreuswieser and Rennenberg 2014, Garssen et al. 2015). 67 In order to cope with the challenges imposed by this scenario, it will be necessary to 68 develop new willow genotypes combining tolerance to flooding with improved growth 69 and wood quality. To evaluate the tolerance of a species to flooding, it is necessary to 70 analyze the responses not only during flooding, but also through the post - flooding 71 recovery period (Striker 2012). For instance, submerged intolerant rice cultivars survive 72 flooding, but suffer from water stress and desiccation upon de - submergence, leading 73 to the death of the plants (Setter et al. 2010). The sudden exposure of previously 74 submerged plants to air may be a stressful situation because of the abrupt raise in  $O_2$ 75 and irradiance, which cause an increase in Reactive Oxygen Species (ROS) or 76 photoinhibition (Luo et al. 2009). Some willow species like Salix variegata develop an 77 increased protection against the post - flooding oxidative damage under complete 78 submergence (Lei et al. 2012).

Apart from the possible damage caused by post - anoxic injury, there are several traits related to productivity in willows that may be affected by flooding, like leaf 81 area, specific leaf area and leaf nitrogen concentration (Robinson et al. 2004, Tharakan 82 et al. 2005). In addition to that, flooding reduces the root - to - shoot ratio in willows 83 (Markus - Michalczyk et al. 2016, Doffo et al. 2017). These morphological and 84 physiological changes are likely to have an impact upon growth during the post -85 flooding period. Willows can be divided into two major ecological groups: riparian 86 species adapted to periodically flooded environments, and wetland species that can 87 grow in lowlands permanently covered with stagnant water (Dickmann and Kuzovkina 88 2014). In this work, we analyzed the progeny of five families, combining parents of S. 89 alba (typically riparian), S. nigra (wetland species), S. humboldtiana (the only native 90 willow species in South America, Dickman and Kuzokvina 2014), and S. matsudana, 91 which is able to endure repeated periods of complete submergence (Wang et al. 2017). 92 Since the parent's habitats experience a variety of flooding regimes, we expected to 93 find different degrees of stress tolerance in the F1 progeny.

The aims of this work were: 1) To analyze the morphological and physiological traits related to productivity in willows during the post - flooding period; and 2) To find out if these traits change differently according to the genotype and the depth of the floodwater. The hypothesis was that the growth recovery in the post - flooding period would be different according to the genotype and the depth of the floodwater.

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100

# 101 Material and Methods

## 102 Plant material, growth conditions and stress treatment

Three genotypes of the F1 of each of five willow crosses were used in this work (15 genotypes in total); the parentage is detailed in Table 1. One family has a typically riparian mother (F9420), three families combine a riparian with a wetland species (F9408, F9802 and F13), and F9813 combines two wetland species. These individuals belong to the breeding program developed by the National Institute of Agricultural 108 Technology (INTA). The genotypes have already passed most selection steps of the109 breeding program, based on their growth, form and pest resistance.

110 One - year - old cuttings of 20 cm long were planted in 3.5 L pots, filled with a 111 1:1 mixture of soil and sand. Before planting, the cuttings were placed in water 112 overnight, and treated with fungicides to avoid diseases. One cutting was planted per 113 pot, and they were placed in a greenhouse with natural irradiance and under natural day length in La Plata (34° 59' 09" S; 57° 59' 42" W). The pots were watered daily, 114 115 keeping the substrate at field capacity. Before the beginning of the treatments, plants 116 were pruned leaving only one shoot per cutting, and fertilized twice with complete 117 Hoagland solution (50 ml per pot, Leggett and Frere 1971).

118 Two flooding experiments were carried out: one with the water level at 10 cm 119 above the soil surface (T10), and a deeper flooding treatment, with the water level at 120 65 cm above the soil surface (T65). In T10, only the root system was flooded, while in 121 T65 most of the shoot was covered by water. The experiments were performed in 122 different years (T10 during 2013 and T65 during 2014); each one had its own set of 123 control plants (watered to field capacity) and differed in duration. Consequently, the 124 statistical analysis was done separately for each of them. A scheme of each 125 experiment is provided in Supplementary Fig.1. The variables measured, their 126 abbreviations and units are detailed in Table 2.

127 For the T10 experiment, the cuttings were planted in pots on August 9, 2013. 128 The treatments were: Control (watered to field capacity), and submerged in water 10 129 cm above soil surface (T10). Flooding started when the plants were 72 days old. The 130 plants were flooded by placing them inside a bigger sealed pot, as previously described 131 (Cerrillo et al. 2013). There were 6 replicates for each genotype and treatment, in a 132 completely randomized layout (N=12 for each genotype, 36 for each family; 18 plants 133 for control and 18 for T10 treatment). The flooding treatment started on October 21, 134 2013 and ended on December 20, 2013. After the end of flooding, a post - flooding 135 recovery period of 30 days started, in which the pots were watered daily to field capacity. The final destructive measurements started on January 20, 2014, marking theend of the experiment.

In the T65 experiment, cuttings were planted on August 13, 2014. The control

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139	plants were watered daily to field capacity and the flooded plants were submerged to
140	65 cm above soil level (T65). The plants in the T65 treatment were placed in a pool
141	filled with water; the water depth in the pool was checked every day and maintained at
142	the same level by replacing the evaporated water when necessary. There were 6
143	replicates for each genotype and treatment, in a completely randomized layout (N=12
144	for each genotype, 36 for each family; 18 plants for control and 18 for T65
145	treatment). The flooding treatment started on October 16, 2014, when the plants were
146	62 days old, and lasted until November 19, 2014. After that date, it followed a post-
147	flooding period until December 15, 2014, when the final destructive sampling started.
148	
149	Growth measurements and leaf traits
150	Height was measured with a ruler, and basal diameter with a digital caliper. The
151	volume index was calculated as follows:
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153 VI= 
$$[(basal diameter)^2 \cdot total height]$$

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155 The Flooding Tolerance Index (FTI, Fichot et al. 2009) was determined using 156 the VI as follows:

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160 The relative growth rates of the stems (RGR), either in height or basal diameter, 161 were determined according to Whitehead and Myerscough (1962). The individual leaf 162 area (ILA) and the specific leaf area (SLA) were determined on the latest expanded 163 leaf at the end of the experiment. The leaf was scanned and the area determined with the software Image J (<u>http://rsbweb.nih.gov/ij/</u>, Schneider et al. 2012). At the end of the experiment, the total biomass for leaves, stem and roots was determined after drying the material at 65 °C to constant weight. Root - to - shoot ratio (RSR) was calculated with those data.

Leaf nitrogen concentration was determined on a pool of leaves, using theKjeldahl method for total nitrogen (Brenmer 1996).

170

## 171 Stomatal conductance and ETR determinations

172 The stomatal conductance (gs) was determined with a Decagon SC1 porometer 173 and the electron transport rate (ETR) with a modulated chlorophyll fluorescence meter 174 (Hansatech FMSII, UK). The measurements were carried out between 10.30 and 13.30 175 h, on cloudless days, on the latest expanded leaf. The average irradiance during the measurements was 967  $\mu$ moles m<sup>-2</sup> s<sup>-1</sup>. Two measurements were carried out in the 176 177 T10 treatment: one during late flooding (53 days after the start of flooding for gs, 54 178 days for ETR) and another during the post - flooding period (24 days after the end of 179 the flooding treatment for ETR, 26 days for gs). For the T65 experiment, 180 measurements were performed one day and 22 days after the end of flooding for ETR, 181 and 9 days and 20 days after the end of flooding for gs.

182

## 183 Statistical Analysis

The statistical analysis was carried out with R 3.2.3 (R Development Core Team 2017), using the package agricolae. The aov function was used for the ANOVA, with clone and treatment as factors, and the post hoc analysis was carried out with the LSD test.

188

189 Results

190 The ANOVA results are depicted in Table 3, showing family, flooding and their 191 interaction as factors. Since T10 and T65 were carried out in different years with their 192 own control treatments, each experiment was analyzed separately.

After one week of flooding, all genotypes developed hypertrophied lenticels andadventitious roots in the submerged parts of the stem (Suppl. Fig.2).

Dry matter accumulation (TDW) and partitioning (RSR) were different in the T10 and T65 treatments (Fig. 1). In T10, TDW was not reduced by flooding, while in T65, it was significantly reduced in all families. The RSR was reduced by both flooding treatments, but the differences were not statistically significant in the T10 treatment for the F13 and F9420 families. In T10, there was a change in dry matter partitioning without total biomass reduction, while in T65 there was a reduction in total biomass plus a change in partitioning.

202 The relative growth rate in height during flooding (RGRh f, Fig. 2) was different 203 in both treatments. In T10 there was no reduction, while in T65, RGRh f was 204 significantly reduced in all families. In the post - flooding period, there were differences 205 in the relative growth rate in height (RGRh pf) according to family and treatment (Fig. 206 2); F9408 increased RGRh pf in both T10 and T65, while F9802 did not. The other 207 families showed different responses according to the treatment, increasing in some 208 cases and without change in others, but there was no significant reduction in RGRh pf 209 in any case.

The relative growth rate in basal diameter during flooding (RGRd f, Fig. 3) in T10 was similar or higher than in control plants, while in T65, it was similar or lower compared to the non - stressed treatment. In the post - flooding period, there were no significant differences in RGRd pf between control and flooded plants except for F9420 in T10.

In the leaves developed during the post - flooding period, there were no
 differences in SLA between control and flooded plants in neither T10 nor T65, but there

were differences among families (Table 3). The size of the leaves developed during the
post - flooding period (ILA) was affected by genotype and treatment (Table 3).

The electron transport rate (ETR, Fig. 4) did not change in T10, neither during flooding nor through the post - flooding period. For T65, no measurements were made during flooding because most leaves were covered by water. One day after the end of flooding, there was an increase in ETR that was statistically significant in three families. This increase did not last in the post - flooding period except for F9813.

Stomatal conductance (gs, Fig. 5) was not affected by the T10 treatment, except for family F9408, which experienced a reduction in the post-flooding period. T65 was measured only in the post - flooding period, and 9 days after the end of the stress episode, there was a significant increase in gs in the previously flooded plants in all families except for F13. This effect did not last long; 22 days after the end of flooding, gs was significantly higher only in F9813.

There were no differences in nitrogen concentration per unit leaf area in the T10 treatment compared to controls (N, Fig. 6), while in T65 it was only significantly increased in F13.

The flooding tolerance index for the volume index (FTI, Fig. 7) was determined at the end of the flooding treatment and again after the post-flooding recovery period. In the T10 treatment, flooded plants had a higher above - ground biomass than controls (FTI higher than 100), while in T65, growth was reduced by flooding (FTI lower than 100). For the T65 treatment, the family ranking was similar after flooding and during the post - flooding recovery period, while in T10 it was different.

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240

241 Discussion

242 Effect of flooding depth on leaf traits related to productivity in willows.

In a previous work, we found that leaf nitrogen concentration increased in deeply
 flooded but not shallowly flooded plants (Rodriguez et al. 2018). These are interesting

245 results, since leaf nitrogen concentration correlates with the photosynthetic rate (Reich 246 et al., 1998) and this could enable a higher photosynthetic fixation rate in the post -247 flooding period. However, we did not find differences in nitrogen concentration between 248 control and T65 plants after 26 days of recovery in four families (the exception being 249 F13). The higher leaf nitrogen concentration did not last long after the end of the 250 flooding episode. The increment occurred in deep flooded willows which experienced a 251 reduction in growth, but not in shallow flooded willows that have a similar biomass as 252 non - flooded plants (Rodriguez et al. 2018). It is possible to speculate that N uptake will 253 continue in flooded willows, as it does in flooded Populus tremula x P. alba plants 254 (Kreuzwieser et al. 2004). Thus, the increase in leaf nitrogen concentration is a 255 consequence of the continuous uptake plus the transient reduction in growth, acting as 256 a reserve that can be used for growth after the end of flooding (Warren et al. 2003).

257 In addition to leaf nitrogen concentration, individual leaf area (ILA) and specific 258 leaf area (SLA) are traits that correlate with productivity in willows (Robinson et al. 259 2004, Tharakan et al. 2005). Both can be modified by flooding: SLA increases in leaves 260 under submergence (Mommer and Visser 2005) and leaf size can be reduced by 261 flooding (Cerrillo et al. 2013). In a previous work, we found that the deeper flooding 262 treatment increased the SLA of leaves expanded during flooding (Rodriguez et al. 263 2018), but we did not find any effect of treatment in the SLA of leaves developed during 264 the post - flooding period. On the other hand, leaf size had only a moderate correlation 265 with dry mass accumulation in T65 (r = 0.49, p < 0.001, n = 150), and a low correlation 266 in T10 (r = 0.27, p≤ 0.01, n = 150).

267

Floodwater depth and genotypes affect growth responses and dry matter partition in
the post - flooding period.

270 The deeper flooding treatment (T65) was a more stressful situation for willows 271 than shallow flooding (T10). Growth in height, and to a lesser extent in diameter, was

272 reduced during flooding in T65, but not in T10. These results were similar to those 273 reported for Salix alba and S. viminalis (Markus - Mychalzcyck et al. 2016) and Alnus 274 japonica (Iwanaga and Yamamoto 2008), where growth was more reduced with an 275 increase in the floodwater level. The restriction on gas exchange imposed by 276 submergence caused a lower rate of carbon fixation that may explain the lower growth 277 in the deep flooding treatment (Luo et al. 2009). However, the occurrence of non-278 stomatal limitations to photosynthesis could not be ruled out. It has been shown that 279 both stomatal and non-stomatal limitations occur in flooded plants of sunflower (Guy 280 and Wample 1984) and poplar (Bèjaoui et al. 2006).

281 The relative growth rates in the post - flooding period were similar or higher in 282 the previously stressed plants compared to the controls. This is probably related to the 283 fact that stomatal conductance and photosynthetic activity (as ETR) during the post -284 flooding period were similar or higher in previously flooded plants compared to the 285 control treatment. In flood - sensitive species, stomatal closure persist beyond the end 286 of the hypoxia (Sojka 1992), but this is not the case for the Salix species analyzed in 287 this work. It seems that willow leaves did not suffer an extensive damage during 288 flooding, allowing for a fast recovery of gas exchange after the end of the stress 289 episode. A similar behavior has been reported for other riparian species adapted to 290 periodically flooded environments (Luo et al. 2009).

T10 and T65 both reduced the root - to - shoot ratio (RSR), because flooding arrest root growth (Jackson and Attwood 1996) and increases root mortality in willows (Markus - Michalczyk et al. 2016, Doffo et al. 2017). The difference between treatments is that T10 combines a reduced RSR with a similar dry matter accumulation as the control treatment, while in T65 there was a reduction in both RSR and total dry weight. In both flooding treatments, RSR still has not reached the same levels as the control plants after the recovery period.

In spite of the recovery of the relative growth rate in previously flooded plants of the T65 treatment, the biomass accumulation was still significantly lower compared to controls after 26 days of recovery, except for family F9813. It is possible that the other families need a longer period to recover to levels similar to those of the control treatment.

303 The responses of growth and leaf variables may be similar in both flooding 304 treatments, but other responses differed among families. The tolerance index to 305 flooding was calculated using volume index, because it showed a good correlation with 306 total dry weight (r = 0.71 for T10 and r = 0.92 for T65). An interesting result is that the 307 tolerance index rating for the families was different at the end of flooding and after the 308 post - flooding period for T10, but it was similar for T65. This is not a major issue for the 309 genotypes used here, since they are all tolerant to T10 conditions. But it is clear that 310 tolerance differs among families for T65, and the genotypes that are more tolerant for 311 T10 will not necessarily behave in the same way with a deeper floodwater level. This 312 should be taken into account to recommend clones to be planted in flood - prone areas. 313 On the other hand, the variation in response of the families analyzed show that it is 314 possible to combine high growth with flooding tolerance in willows, and to select the 315 best willow genotype according to the risk of flooding of the planting site.

316

## 317 Conclusions and perspectives

Our original hypothesis was accepted, since there were differences in the post flooding responses according to the family and the depth of the floodwater. These results highlight the need to evaluate post - flooding responses, and not only the flooding period, in order to improve willow genotypes to be targeted to endure flooding conditions occurring in particular environments. For the deeper flooding conditions, the better performers were the families with *S. matsudana* as mother. These species could be a source of flooding resistance genes to improve willow genotypes destined toareas with risk of deep and prolonged flooding episodes.

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327

# **328** Author Contribution Statement

329 GND, MER and FYO carried out the experimental work. TC selected the genotypes

analyzed and contributed to the experimental design. VMCL designed the experiment,

331 carried out the statistical analysis and wrote the paper.

- 332
- 333

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338

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342

## 343 **Conflict of interest**

344 The authors declare that they have no conflict of interest.

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- 472

473 Table 1 - Plant material used in this work. \*This clone is a spontaneous hybrid between

474 a *Salix humboldtiana* mother and an unknown father.

Family	Mother	Father		
F9408	S. matsudana NZ693	S. alba S7		
F9813	S. matsudana NZ693	S. nigra C7-22		
F9802	S. matsudana NZ692	<i>S. alba</i> SI58-004		
F9420	<i>S. alba</i> SI64-004	Open - pollinated		
F13	S. matsudana	S. x <i>argentinensis</i> cv "Galvete" * x <i>S.</i> <i>alba</i> "114-1"		

478 Table 2 – List of variables measured in this work, with their abbreviations and units.

Variable name	Abbreviations and Units
Root to Shoot Ratio	RSR
Total Dry Weight	TDW (g)
Volume Index	VI (cm <sup>3</sup> )
Relative Growth Rate for height	RGRh (cm day <sup>-1</sup> )
Relative Growth Rate for basal diameter	RGRd (mm day <sup>-1</sup> )
Leaf Nitrogen Concentration	N (μg cm <sup>-2</sup> )
Individual Leaf Area	ILA (cm <sup>2</sup> )
Specific Leaf Area	SLA (cm <sup>2</sup> )
Electron Transport Rate	ETR ( $\mu$ mol electrons m <sup>-2</sup> s <sup>-1</sup> )
Stomatal Conductance	gs (mmol $H_2O \text{ m}^{-2}\text{s}^{-1}$ )

Table 3 – ANOVA table of the variables measured and estimated in this work. The

483 values are those of P. The significant factors (P<0.05) are marked in bold.

Variable		T10			T65	
	Family	Flooding	Interaction	Family	Flooding	Interaction
RSR	0.422	0.0001	0.126	0.0001	0.0001	0.855
TDW	0.0001	0.153	0.871	0.0007	0.0001	0.1444
VI	0.0192	0.0001	0.8106	0.0001	0.0001	0.159
RGRh f	0.0784	0.0169	0.6736	0.1289	0.0001	0.0253
RGRh pf	0.0532	0.0076	0.2906	0.0139	0.0001	0.1450
RGRd f	0.0196	0.0035	0.4969	0.4481	0.0012	0.0549
RGRd pf	0.501	0.490	0.221	0.0477	0.1006	0.9509
Ν	0.771	0.255	0.650	0.0307	0.2683	0.4131
ILA	0.0001	0.0004	0.0838	0.0001	0.0001	0.0355
SLA	0.0001	0.103	0.857	0.0095	0.0867	0.0156
ETR <sup>1</sup>	0.3776	0.6830	0.0818	0.396	0.0001	0.724
ETR <sup>2</sup>	0.113	0.678	0.321	0.358	0.231	0.336
gs <sup>1</sup>	0.0139	0.6414	0.5176	0.0123	0.0001	0.4427
gs <sup>2</sup>	0.0657	0.2035	0.1893	0.0093	0.9096	0.0209

486 <sup>1</sup>: late flooding for T10 and early post-flooding for T65.

487 <sup>2</sup>: post - flooding for T10 and late post-flooding for T65.

489 **LEGENDS TO THE FIGURES** 

490

491 Fig.1 Total Dry Weight (TDW) and Root - to - Shoot Ratio (RSR) in the T10 and T65
492 treatments in five willow families. Means followed by the same letter do not differ
493 according to LSD test (p < 0.05). N=15 for each family and treatment</li>

494

Fig.2 Relative Growth Rate for height (RGRh) for treatments T10 and T65 during flooding (f) and in the post - flooding period (pf) in five willow families. Means followed by the same letter do not differ according to LSD test (p < 0.05). N=18 for each family and treatment

499

**Fig.3** Relative Growth Rate for diameter (RGRd) during flooding (f) and post - flooding (pf) in treatments T10 and T65 in five willow families. Means followed by the same letter do not differ according to LSD test (p < 0.05). N=18 for each family and treatment 503

**Fig.4** Electron Transport Rate (ETR) during flooding and post - flooding for treatment T10 and early and late post - flooding for treatment T65, in five willow families. Means followed by the same letter do not differ according to LSD test (p < 0.05). N=15 for each family and treatment

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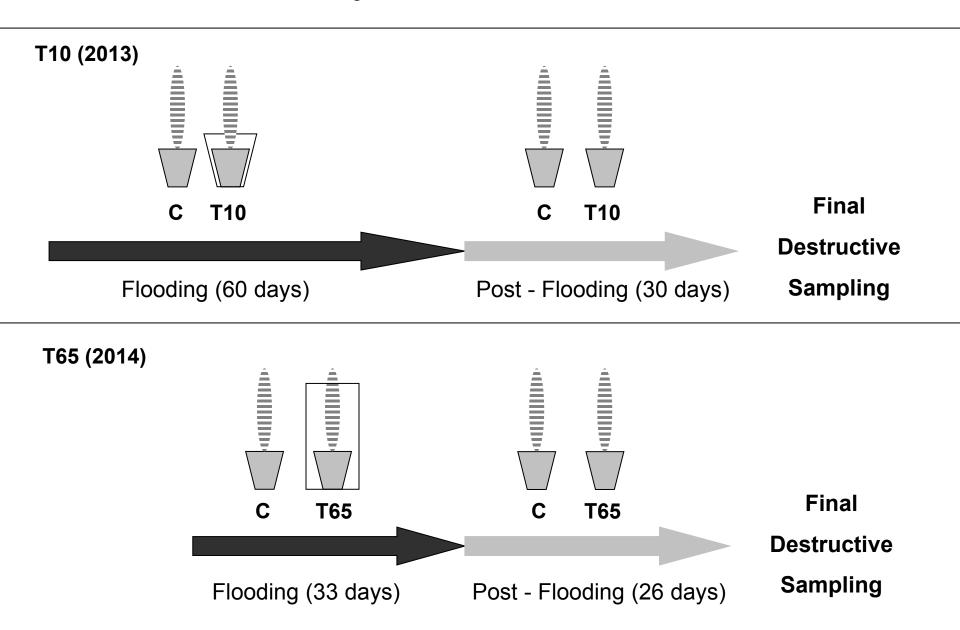
**Fig.5** Stomatal conductance (gs) during flooding and post - flooding for treatment T10 and early and late post - flooding for treatment T65, in five willow families. Means followed by the same letter do not differ according to LSD test (p < 0.05). N=15 for each family and treatment

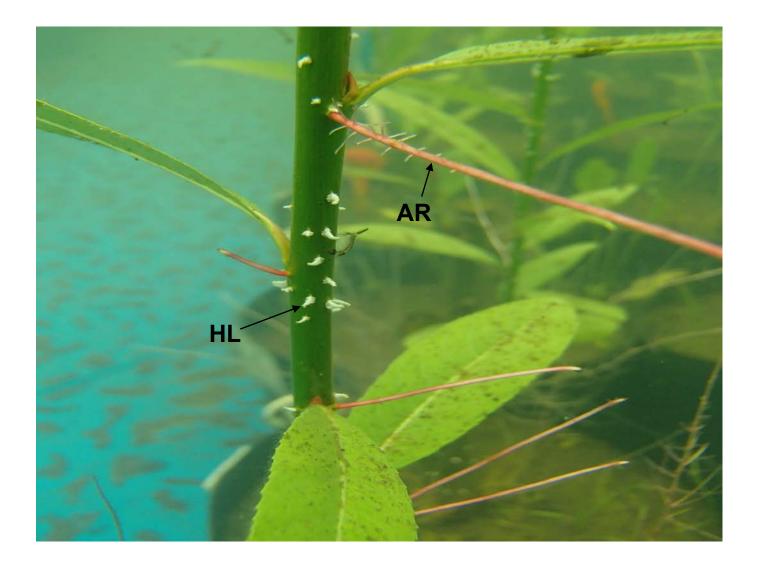
513

**Fig.6** Leaf Nitrogen concentration per unit leaf area (N) at the end of the T10 and T65 experiments, for five willow families. Means followed by the same letter did not differ according to LSD test (p < 0.05). N=12 for each family and treatment 517

**Fig.7** Flooding Tolerance Index (FTI) of the five families, calculated with the Volume Index for both experiments (T10 and T65) at the end of flooding (flooding) and at the end of the post-flooding recovery period (post - flooding). The value was calculated with the average Volume Index for each treatment and family

**Suppl. FIG.1.** An outline of the experiments carried out in this work. T10: Plants submerged 10 cm above soil level. T65: Plants submerged 65 cm above soil level.





**Suppl.FIG.2.** Hypertrophied Lenticels (HL) and Adventitious Roots (AR) developed in the submerged parts of the stems of willow plants.