Changes in durum wheat root and aerial biomass caused by the introduction of the Rht-B1b dwarfing allele and their effects on yield formation

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Abstract
Aims This study aimed to quantify the changes in root and aerial biomass of durum wheat brought about by the introduction of the Rht-B1b dwarfing allele and their effects on yield formation.
Methods A historical series of 24 Mediterranean cultivars with allelic variants a (tall) and b (semi-dwarf) at Rht-B1 locus was tested in tubes in three greenhouse experiments and six field experiments.
Results The dwarfing allele reduced the aerial biomass of each plant at anthesis by 7.6 % and the root by 28.1 % (25.4 %, 26.7 % and 36.0 % in the upper, middle and lower root sections, respectively). Aerial and root biomass were reduced by 27.0 g y⁻¹ and 7 g y⁻¹ respectively, but the relative rate of change was much greater for roots (−0.73 % y⁻¹) than for aerial organs (−0.17 % y⁻¹). Aerial biomass at anthesis was negatively associated with spike number, harvest index and yield in tall cultivars, but no significant relationship was found for semi-dwarf ones.
Conclusions The root/aerial biomass ratio was 29 % lower in semi-dwarf than in tall cultivars. In tall cultivars large aerial biomass at anthesis was detrimental to yield formation, while in semi-dwarf cultivars high aerial biomass at anthesis had no effect on yield formation.

Keywords Breeding effects · Genetic gain · Historical series · PVC tubes · Root section

Introduction

The root system of wheat is essential for firmly anchoring the plant to the soil and taking up water and nutrients from it. In reduced-input agricultural systems, root traits affecting the acquisition of mineral elements often determine yield (Ehdaie et al. 2010; White et al. 2013). In the Mediterranean Basin, one of the largest durum wheat producers in the world, more than 50 % of the total grain of durum wheat is produced in arid and semi-arid conditions, with severe drought most years (Loss and Siddique 1994; Araus et al. 2003; García del Moral et al. 2005). In dryland agricultural systems, a large root system that promotes access to soil water and nutrients is regarded as beneficial for plant growth (Richards 2008), although under terminal drought a greater investment in fine roots at depth would improve yield due to the better access to water and nitrogen (King et al. 2003). Accordingly, root
dry weight at depth has been related to drought adaptation (Lopes and Reynolds 2010).

The introduction of semi-dwarf cultivars with greater resistance to lodging during the Green Revolution in the second half of the 20th century led to major yield gains (Borlaug 2007). Dwarfness in most durum wheat cultivars is controlled by the Rht-B1b (formerly Rht1) allele. In order to incorporate in durum wheat the dwarfing genes already identified and used in bread wheat, crosses between tall durums and semi-dwarf bread wheats were intensively carried out by Dr. Borlaug’s program at CIMMYT from the late 1950’s. The varieties ‘Langdon enano’ and ‘Barrigon Yaqui enano’ (whose pedigree is YAKTANA-54/NORIN-10/BREVOR/3/2*BARRIGON-YAQUI) were the first semi-dwarf durum lines resulting from that crosses (Julio Huerta-Espinio, pers. Comm.) The objective was to transfer the short and stiff straw character in order to allow the intensification of agronomic practices (increased sowing rate, fertilizer applications and water supply). The Rht-B1b allele induces insensitivity to gibberellic acid, thus preventing stem elongation (Peng et al. 1999; Hedden 2003). In addition to the reduced height of the plant, the Rht-B1b allele has well-known pleiotropic effects on plant growth, and morphological and physiological characters (Gale and Youssefian 1985; Li et al. 2011; Rebetzke et al. 2012). It increases yield, mostly under favourable conditions, by altering the proportion of dry matter allocated to grain (De Vita et al. 2007; Royo et al. 2007, 2008; Álvaro et al. 2008a). However, the effect of the Rht-B1b allele on the root system has been poorly explored (Waines and Ehdaie 2007). Previous studies found no consistent association between height genes and root growth and function (Blum 2011). Though the Rht-B1b allele is known to inhibit stem growth, thus making available a surplus of assimilates that are used for thickening the roots (Miralles et al. 1997) without reducing the stem diameter (Calderini et al. 1996), it has been proposed that the enlargement of the root system and its penetration ability might not be under the control of dwarfing genes (Miralles et al. 1997; Kubo et al. 2005). Vigorous shoot growth has been related to vigorous root growth under a wide range of conditions (Mian et al. 1993). However, negative relationships have been identified between root dry weight and plant height in bread wheat (Miralles et al. 1997), and a lower shoot:root ratio has been found in dwarf cultivars (Siddique et al. 1990), seeming to suggest that the selection of cultivars with higher yield and other suitable traits during the 20th century also led to an unintentional selection of those with a reduced investment in root biomass (Zhu and Zhang 2013). Furthermore, modern cultivars containing the Rht-B1b allele have shown an increased capacity to take up water in irrigated environments (Pask and Reynolds 2013), thus demonstrating greater efficiency in the use of water after anthesis than the old tall cultivars (Subira et al. 2015).

Quantifying root characteristics in field experiments is a labour-intensive task because the root crop zone is difficult to access (Motzo et al. 1993; Waines and Ehdaie 2007; Izzi et al. 2008; Botwright Acuña and Wade 2012). Therefore, little attention has been devoted to root systems in breeding programmes in the past, most of which have dealt with the above-ground plant organs (Waines and Ehdaie 2007). A number of root observation methods have been proposed, but due to their cost or slowness for routine screening work in breeding programmes, the most widespread among breeders has been the tube method (Blum 2011). The assessment of the root system is essential because water availability is the most limiting factor for the correct expression of yield potential even in irrigated environments, and projections indicate that less usable water will be available in the future (Pask and Reynolds 2013). The tube method is therefore useful for boosting knowledge of the most appropriate root systems for the drought environments that will be more common in the future.

The effect of breeding activities during the 20th century and the introduction of dwarfing alleles on yield, yield components, biomass production and allocation was previously studied by our team on a historical series of 24 Italian and Spanish durum wheat cultivars released in different periods and with different allelic compositions in the Rht-B1 gene (Royo et al. 2007, 2008; Álvaro et al. 2008a, 2008b, 2008c; Subira et al. 2014). The same set of cultivars was used in the present study with the following aims: (i) to quantify the changes occurring in root biomass due to the introduction of the Rht-B1b dwarfing allele, (ii) to determine the relationship between the changes in root biomass and the changes in aerial biomass, and (iii) to study the relationship between biomass and yield formation in tall and semi-dwarf cultivars. Objectives (i) and (ii) were addressed through experiments in PVC tubes under greenhouse conditions, while objective (iii) was addressed using data of field experiments and relating them to those obtained in the greenhouse experiments.
Materials and methods

Plant material

The plant material used in this study consisted of a historical series of 24 durum wheat (Triticum turgidum L. var. durum) cultivars selected to represent the germplasm grown in Italy and Spain during the last century (Table 1). The set included tall landraces cultivated before 1945, early semi-dwarf cultivars derived from CIMMYT germplasm such as ‘Mexa’, landmark early European cultivars such as ‘Creso’, and cultivars released by local breeding programmes during 1990s in both countries. PCR-based markers specific for the base pair responsible for the semi-dwarf phenotype were used to identify specifically wild-type (Rht-B1a) and mutant (Rht-B1b) allelic variants at the Rht-B1 locus following the methodology described by Ellis et al. (2002), thus allowing classifying the cultivars into two groups: tall, carrying the Rht-B1a allele, and semi-dwarf, with the Rht-B1b allele. This methodology confirmed the presence of the Rht-B1b allele in the genome of the Italian cultivar ‘Adamello’, which had previously been considered not to carry any dwarfing allele (Isidro et al. 2011; Álvaro et al. 2008a, 2008b, 2008c; Royo et al. 2007), according to the results of the test for sensitivity to gibberellic acid (Gale and Gregory 1977).

Experiments in tubes in the greenhouse

Three experiments were conducted under greenhouse conditions during three growing seasons at CIMMYT research station in Toluca, State of Mexico (19°16’N, 99°34’W). Temperatures and daily photoperiod are shown in Fig. 1. Plants were grown in PVC tubes of 120 cm height and 10.2 cm width filled with a soil/sand mixture at a 1:3 ratio and sealed at the bottom with a hole for drainage. Each tube was fertilized with 3.0 g of urea and 0.5 g of triple superphosphate (TSP), and irrigation was provided periodically to prevent water limitation. Experiments consisted of 72 tubes in 2009 and 2010 and 96 in 2012, arranged in randomized complete block designs with three replications in 2009 and 2010 and four in 2012. Planting dates were 2, 10 and 21 December in 2008, 2009 and 2011, respectively. Three seeds of each cultivar (all of them with a diameter between 2.8 mm and 3.5 mm) were sown in each tube and after seedling emergence only two plants per tube were kept.

At anthesis (Zadoks stage 65, Zadoks et al. 1974) plant height was measured from the soil to the top of the spike excluding the awns, and the entire plants, including the roots, were removed from the tubes (Fig. 2). The roots were carefully washed following the methodology described in Blum (2011) and divided into three sections (upper, middle and lower) of equal length. The above-ground biomass and the three root sections obtained

| Tall cultivars (Rht-B1a) | Semi-dwarf cultivars (Rht-B1b) |
|-------------------------|-------------------------------|
| Name | Year of release | Country of origin | Name | Year of release | Country of origin |
| Balilla Falso | <1930 | I | Creso | 1974 | I |
| Razza 208 | <1930 | I | Adamello | 1985 | I |
| Senatore Cappelli | 1930 | I | Simeto | 1988 | I |
| Carlojucci | 1945 | I | Cirillo | 1992 | I |
| Capeiti 8 | 1955 | I | Flavio | 1992 | I |
| Trinakria | 1970 | I | Zenit | 1992 | I |
| Blanco Verdeal | <1930 | S | Camacho | 1975 | S |
| Clarofino | <1930 | S | Esquilache | 1976 | S |
| Pinet | <1930 | S | Mexa | 1980 | S |
| Rubio de Belalcázar | <1930 | S | Ariesol | 1992 | S |
| Bidi 17 | 1950 | S | Senadur | 1995 | S |
| Astigi | 1999 | S | Boabdil | 2000 | S |
Fig. 1 Daily photoperiod (dashed line) and indoor minimum (dotted line) and maximum (solid line) daily temperatures during the three years of greenhouse tube experiments at Toluca’s research station, CIMMYT (Mexico).
from each tube were oven-dried at 70 °C for 48 h and weighed. Dry weight of plant parts was expressed on a single-plant basis.

Field experiments

Six field experiments were conducted with the same set of cultivars in 2003, 2004 and 2005 at two contrasting sites in Spain: Gimenells in the north-east and Chimeneas in the south (Table 2). Experiments followed randomized complete block designs with three replications and plots of 12 m² (8 rows, 0.15 m apart). Sowing rate was adjusted to 400 and 350 fully viable seeds per m² at Gimenells and Chimeneas, respectively. Plots were fertilised following the recommendations for maximizing yields while preventing lodging, and were kept disease- and insect- free with preventive pesticide applications. Anthesis date was recorded for each plot when 50 % of the plants reached this stage and plants within a 50 cm-long row per plot were pulled up at this stage. In the laboratory the number of plants in each sample was recorded and the plants were oven-dried at 70 °C for 48 h to obtain the crop dry weight (CDW, g per m²) and the aerial biomass per plant as the ratio between CDW and the number of plants per m². A second 50-cm-long row was randomly sampled at ripening (Zadoks stage 92) on each plot and used to determine the number of spikes per m² and the number of grains per spike. Harvest index (HI) was obtained on a dry weight basis as the ratio between total grain weight and above-ground biomass of the same sample. The harvest was conducted mechanically at commercial maturity and grain yield was adjusted at 12 % moisture level. Thousand kernel weight was calculated using a sub-sample of the grain obtained.

Statistical analyses

Combined ANOVAs were performed across greenhouse experiments considering the experiment, the
cultivar and their interaction as fixed factors in the model. The sum of squares of the cultivar effect and its interaction were partitioned into differences between alleles at the \textit{Rht-B1} locus and differences within each of them. Means were compared with the Student $t$ test ($P = 0.05$). Absolute (AGG) and relative (RGG) genetic gains were computed as the slope of the linear regression line fitted to the relationship between the absolute or relative value of the trait and the year of cultivar release. Relative values were computed for each cultivar as percentages irrespective of the average value of all cultivars. Linear regression models were fitted to the relationships between: i) traits assessed in field experiments, ii) traits assessed in the greenhouse, and iii) biomass assessed in tubes and yield and biomass related traits determined under field conditions. In all cases mean cultivar data across replications and experiments were used. All analyses were performed with the JMP V.8.0 (SAS Institute Inc. 2009) and Enterprise Guide 4.2 statistical software (SAS Institute Inc. 2006).

### Results

#### Field experiments

Results of the field experiments showing genetic changes in yield, yield components, biomass and HI have been published elsewhere (Royo et al. 2007, 2008; Álvaro et al. 2008a, 2008b). In this study, the examination of the relationships between yield and its components in terms of biomass production and allocation revealed that, for the whole set of cultivars, grain yield under field conditions

#### Table 2  Site and description of the six field experiments conducted in this study

| Site                | Gimenells (North of Spain) | Chimeneas (South of Spain) |
|---------------------|----------------------------|-----------------------------|
| Coordinates         | 41°30’N, 0°20’E           | 37°08’N, 3°49’W            |
| Altitude (m asl)    | 200                        | 684                         |
| Soil texture        | Fine loamy                | Silty clay                  |
| Soil classification | Mesic calcixerolic xerochrept | Loamy calcixerolic xerochrept |
| Harvest year        | 2003 2004 2005             | 2003 2004 2005              |
| Sowing date         | 26 Nov 2002 16 Dec 2003   | 26 Nov 2004 23 Dec 2002 13 Nov 2003 10 Dec 2004 |
| Environmental conditions from sowing to anthesis (SA) |
| $T_m$(°C)           | 8.64 8.13 7.24 10.7 9.81 9.43 |
| ETo (mm)            | 244 206 226 309 290 348   |
| RH (%)              | 80.3 95.6 91.4 68.6 71.2 56.5 |
| WI (rainfall + irrigation, mm) | 170 + 150 202 + 85 62.7 + 150 249 + 40 334 + 40 89.8 + 120 |
| Environmental conditions from anthesis to maturity (AM) |
| $T_m$(°C)           | 17.6 20.0 18.8 21.3 16.2 21.1 |
| ETo (mm)            | 117 140 140 118 139 182   |
| RH (%)              | 78.9 87.9 85.2 45.6 66.9 38.8 |
| WI (rainfall + irrigation, mm) | 62.5 + 0 6.03 + 65 52.9 + 0 0 79.8 + 0 6.00 + 0 |
| Mean yield (kg ha$^{-1}$) | 5378 6495 6670 2571 4327 1425 |

$T_m$: average mean daily temperature; ETo: accumulated reference evapotranspiration computed by the Penman-FAO methodology (Allen et al. 1998); RH: average mean daily relative humidity; WI: water input (rainfall + irrigation).
**Plant Soil (2016) 403:291–304**

![Diagrams and equations](image-url)

**Figure 1:**

- **a)** Graph showing the relationship between aerial biomass per plant (g) and grain yield (kg ha⁻¹) with the equation: \( y = -6.55x + 62.0 \) and \( R^2 = 0.66^{***} \).
- **b)** Graph showing the relationship between aerial biomass per plant (g) and crop dry weight (g/m²) with the equation: \( y = -4.91x + 535 \) and \( R^2 = 0.61^{***} \).
- **c)** Graph showing the relationship between harvest index (%) and aerial biomass per plant (g) with the equation: \( y = 115x + 414 \) and \( R^2 = 0.85^{***} \).
- **d)** Graph showing the relationship between harvest index (%) and grain yield (kg ha⁻¹) with the equation: \( y = -6.55x + 62.0 \) and \( R^2 = 0.86^{***} \).
- **e)** Graph showing the relationship between aerial biomass per plant (g) and number of spikes per m² with the equation: \( y = -49.1x + 535 \) and \( R^2 = 0.61^{***} \).
- **f)** Graph showing the relationship between aerial biomass per plant (g) and number of grains per spike with the equation: \( y = -2.01x + 40.3 \) and \( R^2 = 0.13 \).
- **g)** Graph showing the relationship between aerial biomass per plant (g) and thousand kernel weight (g) with the equation: \( y = 1.07x + 38.5 \) and \( R^2 = 0.05 \).

**Figure 2:**

- **a)** Graph showing the relationship between aerial biomass per plant (g) and grain yield (kg ha⁻¹) with the equation: \( y = -6.55x + 62.0 \) and \( R^2 = 0.66^{***} \).
- **b)** Graph showing the relationship between aerial biomass per plant (g) and crop dry weight (g/m²) with the equation: \( y = -4.91x + 535 \) and \( R^2 = 0.61^{***} \).
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- **g)** Graph showing the relationship between aerial biomass per plant (g) and thousand kernel weight (g) with the equation: \( y = 1.07x + 38.5 \) and \( R^2 = 0.05 \).
was negatively associated with above-ground biomass at anthesis, determined either at the plant level (Fig. 3a) or at the crop level (Fig. 3b). However, when these relationships were assessed for tall and semi-dwarf cultivars separately they were only significant for the former, which showed greater variability for biomass and also for grain yield (Fig. 3a and 3b). Harvest index was significantly and positively associated with grain yield, and accounted for 85 % of yield variations in the model considering all cultivars (Fig. 3c). This relationship was also significant when semi-dwarf and tall cultivars were measured separately, but its reliability was greater for the latter group. In accordance with these results, aerial biomass per plant at anthesis and HI were negatively associated, particularly for the set of tall cultivars, but not for the semi-dwarf ones (Fig. 3d).

The assessment of the relationship between aerial biomass per plant at anthesis and the three main yield components (number of spikes per m², number of grains per spike and grain weight), showed that for the whole set of cultivars the number of spikes per unit area was negatively associated with aerial biomass per plant at anthesis (Fig. 3e), but no significant relationships were observed between the number of grains per spike and grain weight with aerial biomass per plant at anthesis (Fig. 3f and 3g). This pattern was similar for tall cultivars, while for the semi-dwarf cultivars no significant relationship was found between aerial biomass per plant and any of the three yield components.

Greenhouse experiments

The results of the ANOVA showed large significant differences between experiments, but negligible differences between replicates within each of them (Table 3). Differences between cultivars were significant for all the traits assessed and always accounted for more than 21 % of total variation. For plant height the cultivar effect explained 67.1 % of total variation. Differences between cultivars carrying and not carrying the Rht-B1b allele were also significant for all traits, but for plant height they accounted for 54.5 % of total variation. Variability within the semi-dwarf cultivars was greater for all traits than the variability within the tall cultivars, except with regard to plant height.

The introduction of the Rht-B1b dwarfing allele had a greater effect on root biomass per plant, which decreased by 28.1 %, than on aerial plant biomass, which only decreased by 7.6 % (Table 4). Accordingly, the root/aerial biomass per plant ratio was 28.6 % lower in semi-dwarf cultivars than in tall ones. The lower section of the root, the most distant from the root crown, was the one most altered by the Rht-B1b dwarfing allele, as it was reduced by 36 % in the semi-dwarf compared with the tall cultivars, while differences in the upper and middle sections were 25 % and 27 %, respectively (Table 4). The distribution of root weight in the three sections was similar in tall and semi-dwarf cultivars: 52 % and 54 % in the upper section, 26 % and 27 % in the middle section, and 22 % and 19 % in the lower section, respectively (deduced from Table 4).

Among tall cultivars ‘Trinakria’ showed the lowest values for all the traits assessed in the greenhouse, while ‘Blanco Verdeal’ had the largest aerial and total biomass per plant and ‘Bidi 17’ had the largest root biomass per plant and the greatest root/aerial biomass ratio (Supplementary Table 1). The lowest values for root biomass within the semi-dwarf cultivars were recorded for all sections in ‘Simeto’, which also had the lowest root/aerial biomass ratio. On the other hand, the highest values for all traits except plant height and root biomass in the upper section were recorded in the cultivar ‘Camacho’.

Absolute (AGG) and relative (RGG) genetic gains were significant and negative for all traits (Table 5). In absolute terms, the aerial and total biomass per plant decreased at a higher yearly rate than the root biomass. However, in relative terms the decrease was much greater for root biomass (−0.73 g y⁻¹) than for aerial biomass (−0.17 g y⁻¹).

Aerial biomass was positively and significantly associated with root biomass, both for the whole root (Fig. 4a) and for each of its sections (Fig. 4b, 4c and 4d). These relationships were also significant when tall and semi-dwarf cultivars were considered separately, except for the root middle section in tall cultivars (Fig. 4c).

Relationship between traits assessed in tubes

in the greenhouse and yield formation

The analysis of the relationship between aerial biomass per plant assessed in tubes and aerial biomass per plant (Fig. 5a) and per unit area (Fig. 5b) in field experiments showed positive and significant associations in all cases when all cultivars were taken. Moreover, the relationship between aerial biomass per plant in tubes and grain yield (Fig. 5c) showed a similar trend to that observed when both traits were assessed in the field (Fig. 3a). Although aerial biomass per plant in tubes was more than three times that recorded in field experiments, both models showed the same tendency and were equally
significant for the whole set of cultivars. Similarly, the model fitted to the relationship between aerial biomass per plant measured in tubes and HI determined in the field (Fig. 5d) was similar to that obtained when both traits were determined under field conditions (Fig. 3d). The consistency of all these results led us to examine the relationships between root biomass and both yield (Fig. 5e) and HI (Fig. 5f), which, as expected, were negative in both cases, as occurred for aerial biomass.

### Discussion

**Aerial and root biomass in the greenhouse**

The great disparity between the data obtained in the three greenhouse experiments was not unexpected considering the differences in the range of temperatures recorded in them. Previous studies demonstrated great experimental variability in root traits under both conditions.

### Table 3

Percentage of the sum of squares (type III) of the ANOVA for the aboveground and root traits of the 24 durum wheat cultivars of the historical series assessed in three greenhouse experiments

| Source of variation | d.f. | Plant height | Aerial biomass per plant \((a)\) | Root biomass per plant \((b)\) | Total biomass per plant \((a+b)\) | Root/aerial biomass per plant \((b/a)\) |
|---------------------|------|--------------|-------------------------------|------------------------|-------------------------------|----------------------------------|
| Experiment          | 2    | 13.2***      | 30.2***                      | 24.9***                | 30.1***                      | 22.1***                          |
| Rep (Experiment)    | 7    | 0.09         | 1.96                         | 1.53                   | 1.42                         | 1.96                             |
| Cultivar            | 23   | 67.1***      | 21.2***                      | 25.3***                | 22.1***                      | 25.8***                          |
| Between Rht-B1      | 1    | 54.5***      | 10.8***                      | 11.3***                | 11.1***                      | 9.97***                          |
| Within Rht-B1b      | 12   | 5.03***      | 6.02**                       | 11.6***                | 9.81***                      | 6.95***                          |
| Within Rht-B1a      | 10   | 8.14***      | 5.00**                       | 3.05                   | 4.48**                       | 7.29                             |
| Exp x Cultivar      | 46   | 10.7***      | 13.7*                        | 3.64                   | 4.24**                       | 3.14***                          |
| Exp x Between Rht-B1| 2    | 3.12***      | 1.99**                       | 3.77                   | 4.05                          |                                  |
| Exp x Within Rht-B1b| 24   | 3.71***      | 4.30                         | 5.23                   | 4.11                          | 2.75                             |
| Exp x Within Rht-B1a| 20   | 4.16***      | 7.30*                        | 8.99                   | 7.51**                       | 8.19**                           |
| Residual            | 161  | 1.98         | 28.2                         | 25.6                   | 26.9                          | 28.1                             |
| Total               | 239  |              |                              |                        |                               |                                  |

*\(P<0.05; **P<0.01; ***P<0.001\)

### Table 4

Mean values across three greenhouse experiments for the aboveground and root traits of tall (Rht-B1a) and semi-dwarf (Rht-B1b) durum wheat cultivars of the historical series

| Trait                             | Rht-B1a Mean | Rht-B1b Mean | Difference (%) |
|-----------------------------------|--------------|--------------|----------------|
| Mean                               | Range        | Mean         | Range         |               |
| Plant height (cm)                  |              |              |                |
| Aerial biomass per plant \((a)\)  | 88.8b        | 73.5–108     | 60.7 b         | 54.0–75.5     | −31.6          |
| Total root biomass per plant \((b)\) | 17.0 a       | 15.4–17.9    | 15.7 b         | 15.1–17.2     | −7.60          |
| Root biomass per plant upper section (g) | 1.14 a       | 0.85–1.33    | 0.82 b         | 0.56–1.28     | −28.1          |
| Root biomass per plant middle section (g) | 0.59 a       | 0.43–0.77    | 0.44 b         | 0.31–0.66     | −25.4          |
| Root biomass per plant lower section (g) | 0.30 a       | 0.25–0.37    | 0.22 b         | 0.15–0.34     | −26.7          |
| Total biomass per plant \((a+b)\)   | 18.1 a       | 16.3–19.2    | 16.6 b         | 15.6–18.5     | −8.29          |
| Root/aerial biomass per plant \((b/a)\) | 0.07 a       | 0.055–0.079  | 0.05 b         | 0.036–0.074   | −28.6          |

Means within rows followed by the same letter are not significantly different according to the Student’s t test at \(P=0.05\)
Table 5 Absolute (AGG) and relative (RGG, % y⁻¹) genetic changes for the aboveground and root traits of the 24 durum wheat cultivars of the historical series assessed in three greenhouse experiments

| Trait                                         | R²    | AGG  | RGG  | AGG units |
|-----------------------------------------------|-------|------|------|-----------|
| Aerial biomass per plant (g) (a)              | 0.66***| -27.0 | -0.17 | g · y⁻¹   |
| Total root biomass per plant (g) (b)          | 0.54***| -6.99 | -0.73 | g · y⁻¹   |
| Root upper section biomass per plant (g)      | 0.47***| -3.42 | -0.67 | g · y⁻¹   |
| Root middle section biomass per plant (g)     | 0.51***| -1.68 | -0.67 | g · y⁻¹   |
| Root lower section biomass per plant (g)      | 0.61***| -1.83 | -0.90 | g · y⁻¹   |
| Total biomass per plant (g) (a + b)           | 0.65***| -34.0 | -0.20 | g · y⁻¹   |
| Root/aerial biomass per plant (b/a)           | 0.48***| -0.32 | -0.56 | y⁻¹       |

***p < 0.001

greenhouse conditions (Botwright Acuña et al. 2007) and field conditions (Botwright Acuña and Wade 2012). However, the results of the three replicates of each experiment were consistent, as shown by the lack of statistical significance for all traits of the replicate effect (nested to the experiment) of the ANOVA.

![Graphs showing the relationship between traits assessed for tubes in the greenhouse.](image)

**Fig. 4** Relationship between traits assessed for tubes in the greenhouse. Each point corresponds to the mean data of a durum wheat cultivar across three experiments and three (2009 and 2010) or four (in 2012) replications per experiment. Relationships between aerial biomass per plant and a total root biomass, b upper 1/3 root section biomass, c middle 1/3 root section biomass, and d lower 1/3 root section biomass. Black circle Tall cultivars (Rht-B1a), White circle Semi-dwarf cultivars (Rht-B1b). Regression equations refer to the models fitted to all cultivars (n = 24). Coefficients of determination of linear models fitted to tall (n = 11) and semi-dwarf (n = 13) cultivars are also shown on each figure. **p < 0.01; ***p < 0.001
significance of the cultivar effect for all traits and the high percentage of total variability explained by its sum of squares denoted the wide genetic diversity in the set of cultivars used in the current study. Particularly for plant height, the cultivar effect accounted for ca. 67 % of total variation, and differences between Rht-B1 alleles explained more than 81 % of genotypic variability, thus supporting the importance of plant height in differentiating durum cultivars released before and after the Green Revolution (Royo et al. 2007, 2008; Graybosch et al. 2006).

**Fig. 5** Relationships between traits assessed for tubes in the greenhouse (x axis) and traits assessed in the field (y axis). Relationships between aerial biomass per plant in tubes and a aerial biomass, b crop dry weight, c yield and d harvest index. Relationships between root biomass per plant and e yield and f harvest index. Each point corresponds to one of 24 durum wheat cultivars. Regression equations refer to the models fitted to all cultivars (n = 24). Coefficients of determination of linear models fitted to tall (n = 11) and semi-dwarf (n = 13) cultivars are also shown on each figure. ***P < 0.001
and Peterson 2010). The greater variability of aerial and root biomass found within semi-dwarf cultivars when compared with tall ones supports the statement that genetic variability was not reduced in modern Mediterranean durum wheat cultivars as a consequence of the breeding activities conducted during the 20th century (Martos et al. 2005).

The significant differences observed in the greenhouse between cultivars carrying the Rht-B1a (tall) and Rht-B1b (semi-dwarf) allele for aerial and root biomass indicated that the dwarfing allele not only reduced above-ground biomass, as widely reported by previous studies (Brancourt-Hulmel et al. 2003; Royo et al. 2007; Álvaro et al. 2008a), but also caused a decrease in total root biomass and in each root section. Moreover, the reduction caused by the dwarfing allele in root biomass (ca. 28 %) was much greater than that observed in aerial biomass (ca. 8 %), as shown by the relative rate of change, which was more than four times higher for the roots than for the aerial organs (−0.73 % y⁻¹ and −0.17 % y⁻¹, respectively). Nevertheless, as the aerial fraction of the plant was much heavier than the root fraction, when expressed in absolute terms the genetic change was about four times greater for the above-ground biomass (−27.0 g y⁻¹) than for the roots (−7 g y⁻¹). The relative change in the root/aerial biomass ratio was more than twice that recorded for aerial or total biomass, showing that in relative terms the dwarfing allele had a greater effect on reducing the dry matter of roots than on reducing that of aerial organs. The root/aerial biomass ratio was 29 % lower in semi-dwarf cultivars than in tall ones. Considering the higher grain yield of the former (Canevara et al. 1994; De Vita et al. 2007; Motzo et al. 2004; Royo et al. 2008), this finding may suggest that cultivars carrying the Rht-B1b allele have greater root efficiency for input capture than those carrying the Rht-B1a allele, which is in agreement with the recent proposal of Subira et al. (2015) that semi-dwarf cultivars have an improved capacity to respond to water availability after flowering. In addition, the larger biomass at depth of tall cultivars may denote greater transpiration under drought stress (Blum 2011).

The strong relationship found between root and aerial biomass assessed in tubes was consistent with the reduction caused in both of them by the Rht-B1b dwarfing allele. However, the slope of the linear regression model fitted to the relationship between aerial biomass and biomass of the lower root-section was more than twice the slope of the model fitted to the upper root-section. This result may indicate that differences between cultivars in root biomass are related to greater differences in aerial biomass when the divergence is due to the lower rather than the upper part of the root.

Relationship between aerial biomass and yield in the field

The results of the study of the relationships between aerial biomass and grain yield under field conditions depended on the specific set of cultivars that were considered when the models were fitted. For the whole set of cultivars, around 60 % of yield variations were explained by differences in the aerial biomass, considered either on a plant or a crop basis, which were negatively associated with yield. Moreover, for the whole set of cultivars 85 % of yield variations were explained by genotypic differences in HI, which had a positive relationship with yield and a negative association with aerial biomass. However, when semi-dwarf and tall cultivars were analysed separately, these relationships were statistically significant only for tall cultivars, suggesting that for this sub-set large biomass at anthesis resulted in low yields, probably due to its negative association with HI. These results indicate that, for the set of cultivars carrying the Rht-B1a allele used in the current study, the maintenance of a large crop biomass competed with the allocation of dry matter in the grain or, in other words, large biomass at anthesis was detrimental for achieving high yields. In contrast, for the set of semi-dwarf cultivars, yield did not depend on the crop biomass at anthesis but was positively associated with HI. These results suggest that, for the set of cultivars carrying the Rht-B1b studied here, yield relied on the capacity of the plant to accumulate photosynthates in the grain, independently of the size of the plant canopy at anthesis. It has been demonstrated that one of the greatest effects of the dwarfing allele was the increase in HI, related to the increase in the number of grains per spike and spikelet and an improved translocation efficiency of pre-anthesis assimilates to grains (Royo et al. 2007, 2008; Álvaro et al. 2008b, 2008c).

Differences between tall and semi-dwarf cultivars were also detected when the relationships between aerial biomass and the three main yield components were examined in field experiments. Although both number of grains per spike and grain weight were independent of the plant biomass at anthesis regardless of the set of cultivars considered, for genotypes carrying the Rht-B1a...
alleles the number of spikes per unit area was negatively associated with aerial biomass, while the two traits were independent for semi-dwarf cultivars. These results may suggest competition in tall cultivars between the investment of resources in enlarging the canopy and the development of reproductive organs. This competition did not seem to exist within semi-dwarf cultivars, probably because their lower plant biomass allowed them to develop a larger number of spikes, as reported by Royo et al. (2007).

Aerial biomass assessed in tubes was positively and significantly associated with that obtained in the field when all cultivars were included in the model, but not for each set independently. These results demonstrate that, although plant development in tubes was much greater than that recorded under field conditions at commercial plant densities, the results obtained in tubes and in the field were properly correlated, but only when the model included a wide range of variability. This assumption was confirmed when similar trends were found in the relationships between yield and aerial biomass assessed in both the field and in tubes, and similar patterns were also obtained for the relationships between HI and aerial biomass determined by field and tube experiments when all cultivars were included in the model. In order to predict the relationship between root biomass and yield, it was hypothesized that if a significant association existed between the aerial biomass assessed in tubes and in the field for the whole set of cultivars, a relationship would also most likely exist between root biomass in tubes and in field experiments. In accordance with this assumption, the negative associations that appeared between root biomass and both yield and HI when all cultivars were considered in the model confirm the findings of previous studies regarding the lower efficiency of the biomass of tall varieties in terms of yield formation (Álvaro et al. 2008a, 2008c).

Conclusions

The introduction of the Rht-B1b allele in modern durum wheat cultivars resulted in a reduction in aerial biomass of the plant and an even greater reduction in root biomass along its whole length. A strong and significant positive relationship was found between the aerial and root biomass in the three sections, thus showing that the reduction of the root biomass due to the introduction of the Rht-B1b allele occurred in both the upper and the lower parts of the root. Within the tall cultivars, those with large aerial biomass had a lower number of spikes per unit area and also a lower HI, which resulted in yield reductions. On the other hand, within the semi-dwarf cultivars a greater aerial biomass was neither detrimental nor beneficial in terms of yield formation.

Acknowledgments J. Subira was the recipient of a PhD grant from INIA, Spain. This study was partially funded by projects AGL2002-04285, AGL-2012-37217 and RTA2012-010. Authors thank Dolores Villegas and Julio Isidro for their contribution to the development of experiments.

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