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## Introduction

With harvest index approaching a plateau in some regions and countries, future genetic improvement in grain yield of wheat (*T. aestivum* L.) will increasingly depend on raising above-ground biomass. Therefore, a key breeding target is enhancing photosynthetic capacity to boost biomass. In bread wheat, there are to date relatively few reports of genetic variation in light-saturated leaf photosynthetic rate ( $A_{max}$ ) in field investigations within adapted elite germplasm (Richards, 2000; Foulkes *et al.*, 2009). This study therefore aims to identify novel variation for photosynthetic capacity and biomass by screening diverse wheat germplasm (landraces and synthetic-derived hexaploids) for flag-leaf photosynthetic rate and associated traits and biomass.

## Materials and methods

Field experiments were carried out in 2010-11 and 2011-12 at University of Nottingham farm, Leicestershire UK (52°50' N, 1°14' W) on 50 wheat genotypes (20 wheat landraces from the AE Watkins collection, 20 synthetic-derived hexaploid wheats in a spring wheat Paragon background (BC1S3) and 10 UK elite wheat cultivars). The experiments were grown with optimal inputs and there were 3 replicates. Genotypes in the present analysis were selected to have similar anthesis dates. Flag-leaf  $A_{max}$  was measured using a Li-Cor 6400XT Photosynthesis System weekly from GS39 to mid grain fill on three flag leaves per plot in a subset of 15 genotypes. Grain yield, harvest biomass and relative chlorophyll content (SPAD) at anthesis on three flag leaves per plot were recorded in 50 genotypes.

## Results

Significant genetic variation in biomass ( $P < 0.001$ ) was found in the 50 genotypes with higher biomass overall recorded in the synthetic derivative (1874 g m<sup>-2</sup>) and elite UK cultivar (1756 g m<sup>-2</sup>) groups compared to the Watkins landrace group (1543 g m<sup>-2</sup>) ( $P < 0.05$ ). Genotypes differed in  $A_{max}$  ( $P < 0.001$ ) averaged across readings and years with synthetic hexaploid wheat (22.3-25.0  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ) and elite cultivars (23.2-25.3  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ) generally expressing higher  $A_{max}$  than the Watkins landrace group (18.0-21.5  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ). The year x genotype effect was not statistically significant.

Flag-leaf  $A_{max}$  was positively associated with biomass at harvest (pre-anthesis  $A_{max}$  ( $R^2 = 0.80$ ,  $P < 0.001$ ); post-anthesis  $A_{max}$  ( $R^2 = 0.61$ ,  $P < 0.001$ ) Fig. 1). Similarly, flag-leaf  $A_{max}$  pre-anthesis ( $R^2 = 0.87$ ,  $P < 0.001$ ) and post-anthesis ( $R^2 = 0.74$ ,  $P < 0.001$ ) was positively associated with grain yield. A significant and positive correlation was observed between biomass and grain yield (Fig. 2). A positive association was also found between flag-leaf SPAD value at GS61 and pre-anthesis  $A_{max}$  ( $R^2 = 0.82$ ;  $P < 0.001$ , Fig. 3).

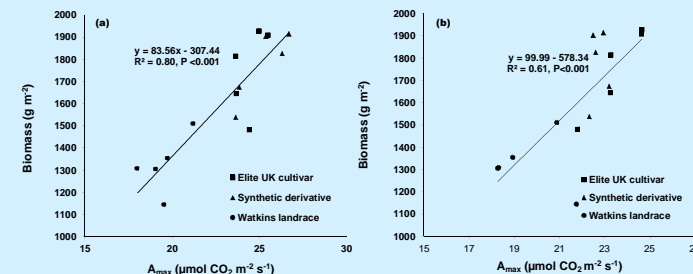


Figure 1: Relationship between pre-anthesis (a) and post-anthesis (b) flag-leaf  $A_{max}$  and biomass (Mean 2010-11 and 2011-12). LSD (5%) for biomass = 562 g m<sup>-2</sup>.



Field experiment at University of Nottingham Farm in 2011-12.

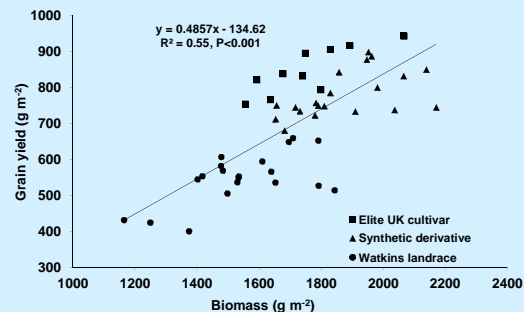


Figure 2: Relationship between grain yield and biomass (Mean 2010-11 and 2011-12).

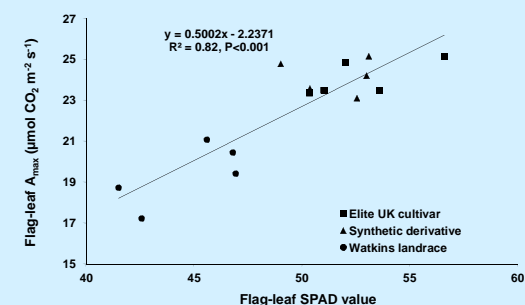


Figure 3: Relationship between pre-anthesis  $A_{max}$  and chlorophyll content (SPAD) (Mean 2010-11 and 2011-12). LSD (5%) for SPAD = 4.81.

## Conclusions

Novel genetic variation in biomass was identified amongst groups of synthetic-derived wheats (1653-2170 g m<sup>-2</sup>), UK elite cultivars (1555-2066 g m<sup>-2</sup>) and AE Watkins landraces (1163-1842 g m<sup>-2</sup>). Genetic variation in flag leaf  $A_{max}$  in the range 18.0 – 25.3 mol  $\mu\text{CO}_2 \text{ m}^{-2} \text{ s}^{-1}$  was positively associated with biomass and grain yield amongst the genotypes. The elite UK cultivars and synthetic-derived wheats generally showed higher  $A_{max}$  compared to AE Watkins landraces. In the present study results showed a linear relationship between flag-leaf N concentration (as indicated by SPAD) and  $A_{max}$  amongst genotypes, in contrast to previous reports of an asymptotic relationship, e.g. Evans (1983). Work is ongoing screening a wider range of synthetic derivatives and landraces for biomass and underlying traits. The strong correlation between  $A_{max}$  and SPAD indicates that SPAD could be used as a selection criteria for high throughput phenotyping.

## References

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