

## **TBD**

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## 2 ABSTRACT

- 3 Hello, this is the abstract.
- 4 Keywords: keyword1, kw2, kw3... have to have at least 5, max 8

## 1 RESULTS

5 ... a quantitative comparison of the datasets in Table 1.

Sample et al. **Running Title** 

**Table 1.** Quantitative metrics of the generated and existing functional annotation sets. C, F, P, and A refer to the aspects of the GO: Cellular Component, Biological Function, Molecular Process, and Any/All.

-			Annotations <sup>a</sup>				Annotated Genes <sup>b</sup>				Median Ann. per G.c			
Genome	Genes	Dataset	C	F	P	A	C	F	P	A	$\overline{\mathbf{C}}$	F	P	A
maize_v3	300	GOMAP Gramene49 Phytozome	135251 20072 4787	87953 31139 19098	291855 30102 13100	515059 81313 36985	34867 11834 4524	38099 18033 13775	39469 15800 11365	39469 21926 16132	3 1 0	2 1 1	6 1 1	11 3 2
maize_v4	320	GOMAP	88831	82849	278952	450632	36717	37431	39324	39324	2	2	6	10
rice	200	GOMAP	72782	64783	248713	386278	28619	29876	35824	35825	2	2	6	9
soybean	52872	GOMAP	129215	114010	417575	660800	46020	47087	52871	52872	2	2	6	11
wheat	500	GOMAP	267742	218839	786028	1272609	95604	98224	107890	107891	2	2	6	10

<sup>&</sup>lt;sup>a</sup> How many annotations in the C, F, and P aspect does this dataset contain? A = How many in total? A = C + F + P

b How many genes in the genome have at least one GO term from the C, F, P aspect annotated to them? A = How many at least one from any aspect?  $(A = C \cup F \cup P)$ <sup>c</sup> Take a typical gene that is present in the annotation set. How many annotations does it have in each aspect? A = How many in total? Ask your favorite statistician why  $A \neq C + F + P$