

IWGSC Phase II: What's Next for the IWGSC

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IWGSC Standards & Protocols Workshop
Plant & Animal Genome Conference

San Diego, California
16 January 2018



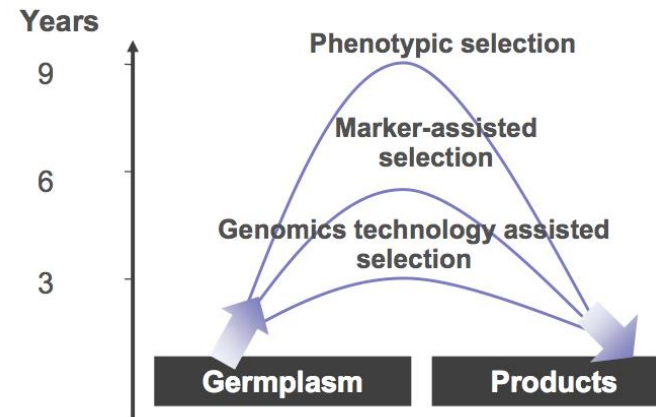
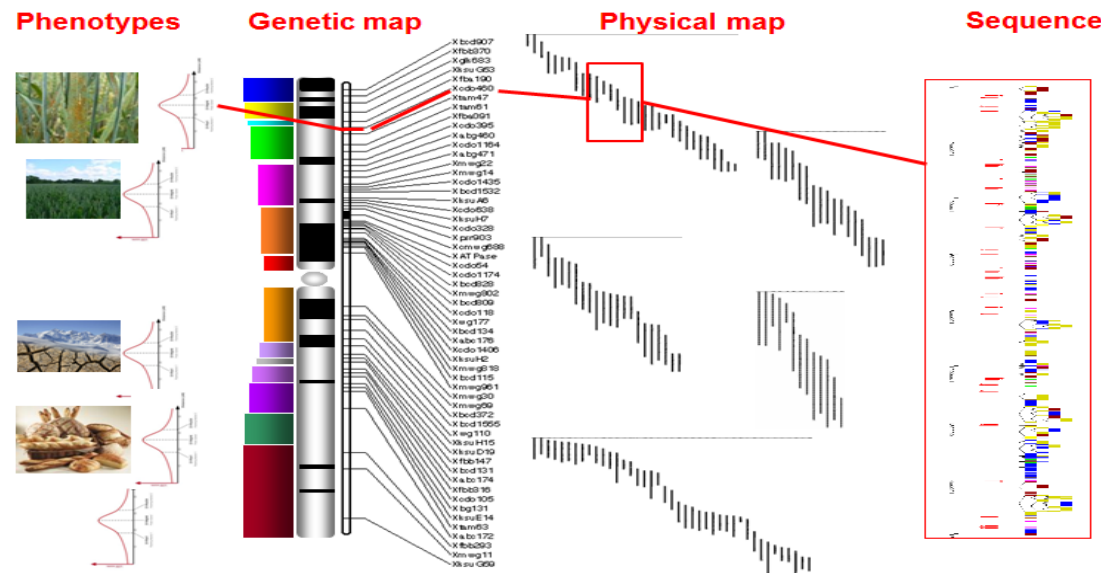
2005 – IWGSC Goal & Vision

Goal

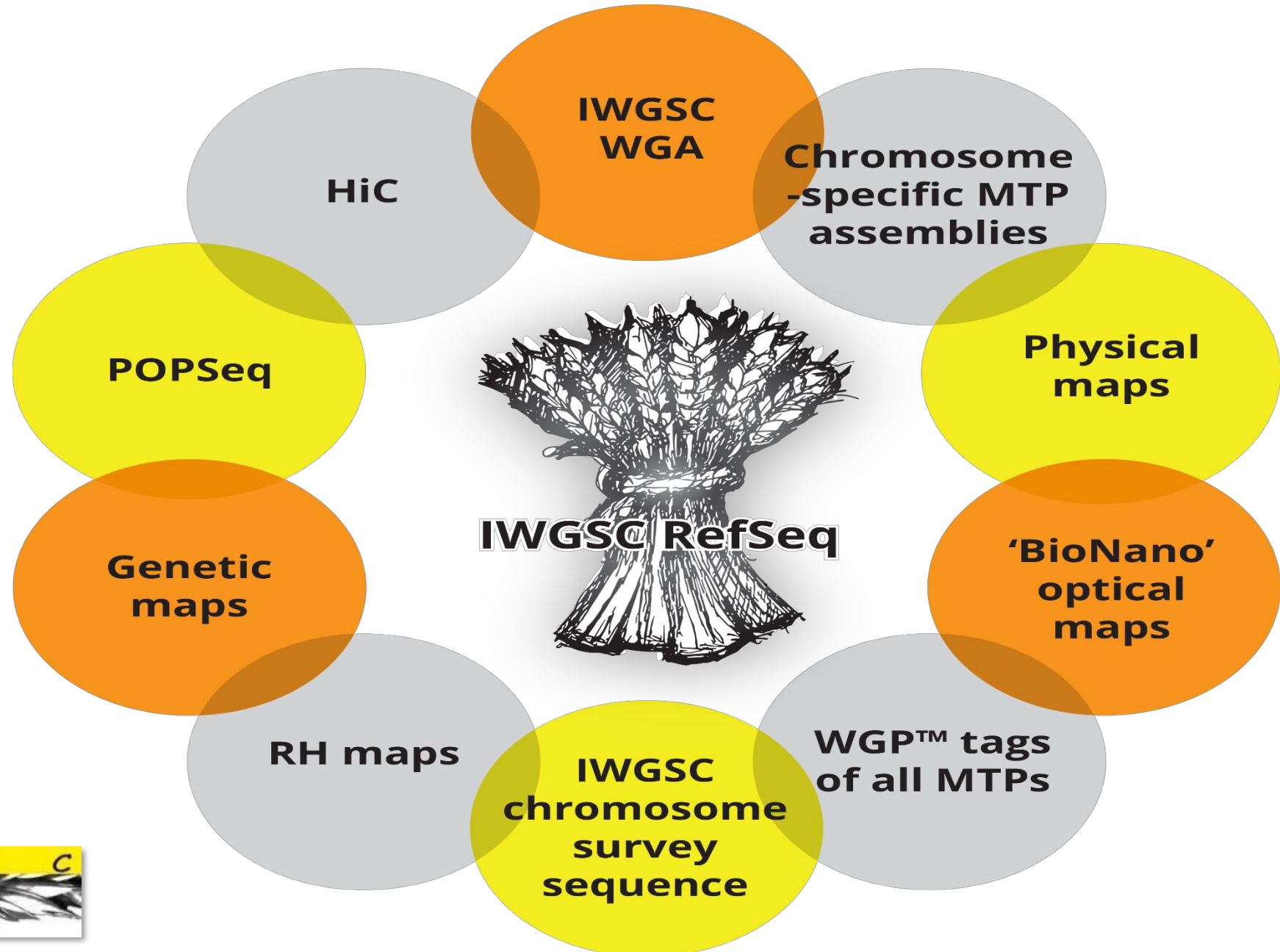
- Lay a foundation to accelerate wheat improvement
- Increase profitability throughout the industry

Vision

- High quality annotated genome sequence, comparable to rice
- Physical map-based, integrated and ordered sequence



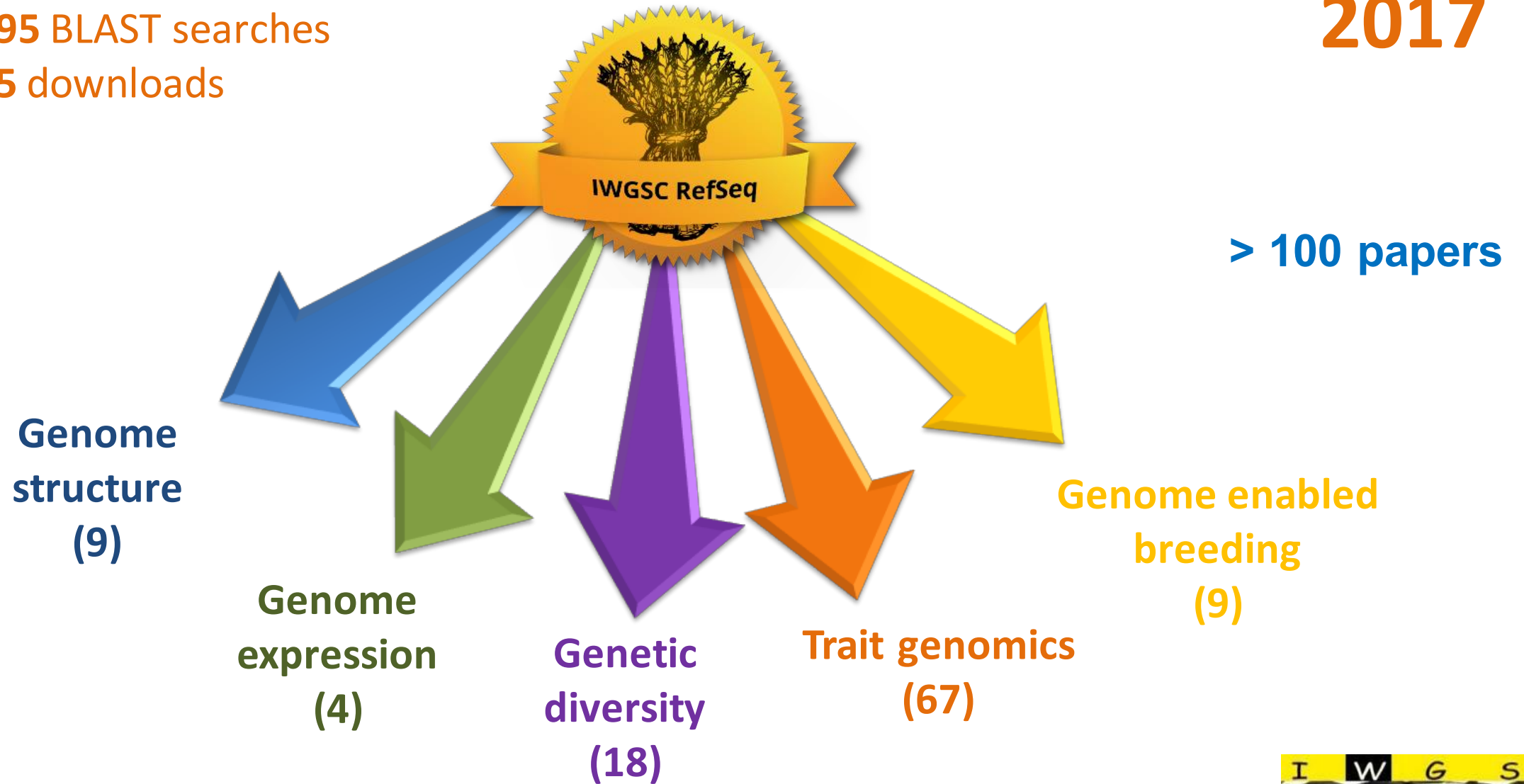
Concerted integration of resources: RefSeq v1.0



Measuring the Impact of IWGSC RefSeq v1.0

476,595 BLAST searches
22,935 downloads

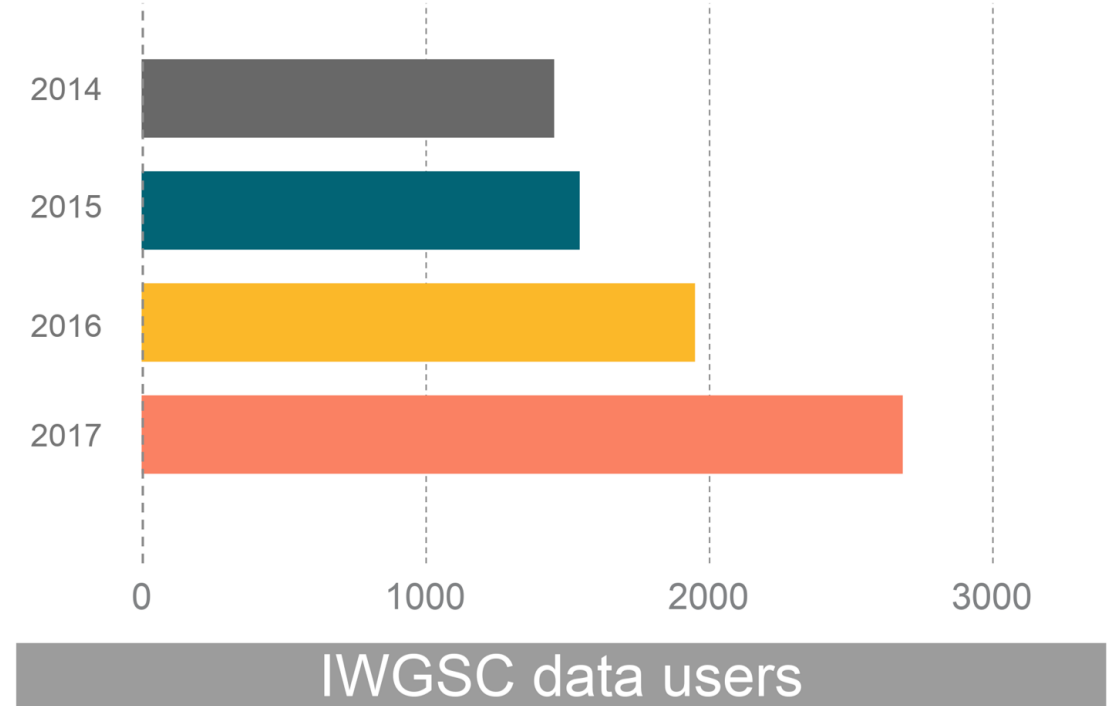
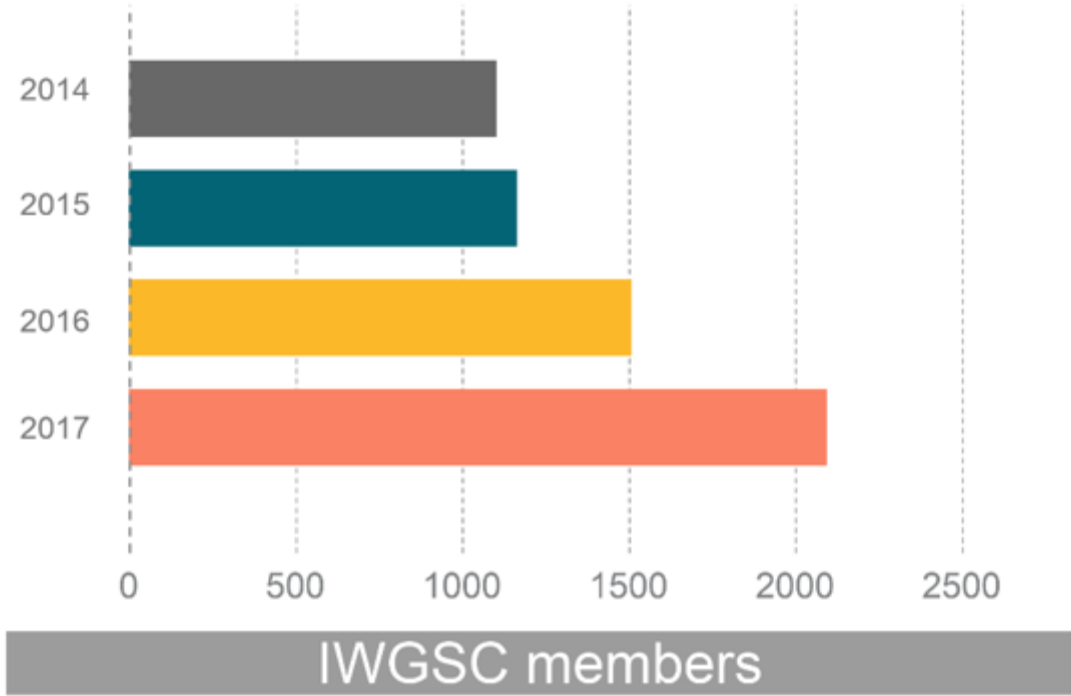
2017



of papers referencing use of IWGSC RefSeq related resources since January 2017



Measuring the Impact of IWGSC RefSeq v1.0



Progress

2005
Vision

- High quality annotated genome sequence, comparable to rice
- Physical map-based, integrated and ordered sequence



Looking into the Future

- **Achieve Gold Standard Reference**
- **Reach beyond reference to ensure a full genome-sequence based toolbox for wheat**



Goal: Accelerate wheat improvement

Vision and strategy for the IWGSC 2.0

Reference sequences with high quality functional annotations

Enhance breeding through an increased understanding of molecular basis of traits and their allelic diversity

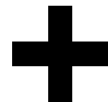
A wheat Pan-genome representing the worldwide diversity

An international, well connected and coordinated community

Public, user-friendly, integrated databases and tools for all users



From RefSeq v1.0 to IWGSC Gold Standard



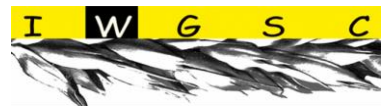
Manual &
Functional
Annotation



- Pipeline for community generated manual annotation with future annotation releases
- Functional annotation
 - Gene family leaders
 - Applying machine learning techniques
- Closing gaps by incorporating new resources to build a IWGSC RefSeq v2.0
- IWGSC leaders: Rudi Appels, Fred Choulet.....

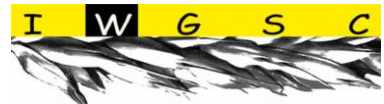
IWGSC 2.0 Activities: Pan-Genome

- Develop a wheat pan-genome that represents the breadth of worldwide wheat diversity
- Reference quality for 8-12 landraces
- Diversity panels at draft quality
- Haplotype database
- Skim-sequencing and high throughput genotyping of many lines
- IWGSC team: Etienne Paux, Sébastien Praud,.....



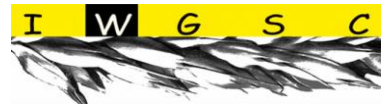
IWGSC 2.0 Activities: Database

- Develop user-friendly, integrated databases and tools
- IWGSC Pan-genome Database Task Force
 - Fred van Ex, Bayer CropScience
 - Magalie Leveugle, Biogemma
 - Matthieu Conte, Syngenta
 - Michael Alaux, URGI-INRA
 -
- What tools do you want?



IWGSC 2.0 Activities: IWGSC Exome Array

- Wheat exome capture array based on IWGSC RefSeq annotation v1.1.
- Accepting data until 31 January 2018 (see- Burkhard Steuernagel)
- Arbor Biosciences will offer enrichment reagents and an enrichment service
- Exome Team: Burkhard Steuernagel, Sreya Ghosh, Sébastien Praud, Hikmet Budak, Etienne Paux.....
- Discussion: What modules would you like to see?



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Opportunity for Community Input

Thank you for your attention!