# Scalable and Diverse

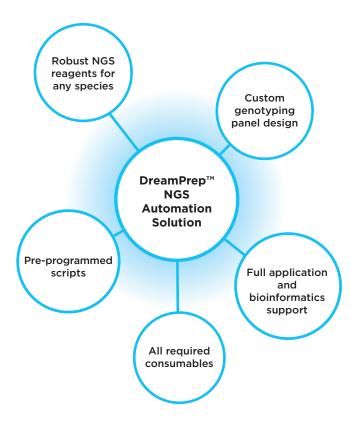
## Solutions for Agrigenomics.

#### Trait to table: Solutions to advance agrigenomics

Agricultural researchers are increasingly adopting next-generation sequencing (NGS) technology as an essential research and development tool in plant and animal genomics. NGS is being used for marker-assisted selection (MAS) to accelerate plant breeding and selection, as well as in transcriptomics, plant or animal-pathogen interactions and epigenetics.

#### Agrigenomics made easy

At Tecan, we work to tailor the right scalable solution to your NGS needs— combining NGS reagents with automation platforms to maximize your productivity.



#### Automation-ready kits for various applications

#### Genotyping by Sequencing

#### Allegro<sup>\*</sup> Targeted Genotyping V2

Fast, scalable, cost-effective editable content for targeted GBS with high sample multiplexing capabilities (100,000+ SNPs)

#### **Microbiome Analysis/Viral Surveillance**

**Allegro' Targeted Genotyping V2** Custom panel design for profiling microbial communities

#### Revelo<sup>™</sup> RNA-Seq High Sensitivity

Effective and high sensitive detection of viral infections in challenging livestock samples, even samples with degraded RNA (e.g., environmental, livestock)

#### Whole Transcriptomics

Universal Plus<sup>™</sup> Total RNA-Seq

End-to-end solution combining library preparation, customizable post-library targeted depletion, and library quantification

#### **Ultra-low Input**

#### Ultralow<sup>™</sup> V2 DNA-Seq

Simple and efficient workflow for ultra-low input as low as 10 pg with high-fidelity amplification across a broad range of GC content

#### High-resolution genotyping meets high throughput and flexibility

Allegro Targeted Genotyping enables highly accurate SNP detection with very high reproducibility and low cost per data point.

- Enable large scale genotyping | Dual-index sample barcoding enables multiplex sequencing of 3000+ samples in a single sequencing lane.
- Automate your workflow | Allegro on DreamPrep NGS is a high-throughput genotyping solution capable of processing up to 384 samples in one day with no user intervention.
- **Target SNPs of interest** | Custom genotyping panels are available for any sequenced species with as many as 100,000 SNPs.

We provide bioinformatics support to generate optimal probe design. A single probe per locus can be used to maximize the number of genotyped sites, while a two-probes design can be leveraged to maximize coverage uniformity and additional resilience to the assay. Sequencing coverage is shown for a small subsample of sequenced reads.

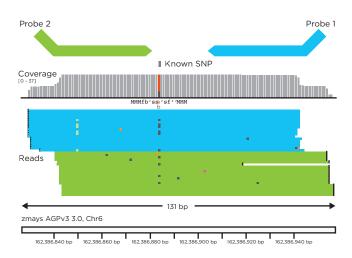


Figure 1: **Very high accuracy genotyping.** Zea mays genotyping panel showing a two-probe design. Probes are designed on both sides of the SNP within 100 bp. Two independent probes interrogate every SNP, enabling very high accuracy SNP-based genotyping.



#### ALLEGRO TARGETED GENOTYPING V2 ON DREAMPREP NGS

Figure 2: **Allegro Targeted Genotyping V2 on DreamPrep NGS.** Our complete targeted genotyping by sequencing solution enables hands-free library preparation for the labor-intensive first steps of the protocol, starting with up to 384 samples and ending with 8 pooled samples ready for the final hybridization and amplification steps.

#### www.tecan.com

#### For Research Use Only. Not for use in diagnostic procedures.

Australia +61 3 9647 4100 Austria +43 62 46 89 330 Belgium +32 15 42 13 19 China +86 21 220 63 206 France +33 4 72 76 04 80 Germany +49 79 51 94 170 Italy +39 02 92 44 790 Japan +81 44 556 73 11 Netherlands +31 18 34 48 17 4 Nordic +46 8 750 39 40 Singapore +65 644 41 886 Spain +34 93 595 25 31 Switzerland +41 44 922 89 22 UK +44 118 9300 300 USA +1 919 361 5200 Other countries +41 44 922 81 11

Tecan, Allegro, Universal Plus, Ultralow, Revelo and DreamPrep are registered trademarks and trademarks of Tecan Group Ltd., Männedorf, Switzerland or of Tecan Genomics, Inc., Redwood City, USA.

s 2022 Tecan Genomics, Inc., all rights reserved. For disclaimer and trademarks please visit www.tecan.com

### ..............................

