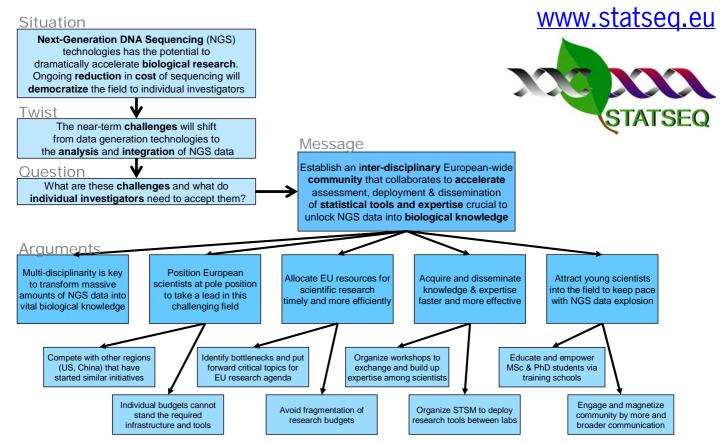




Statistical challenges on the 1000€ genome sequences in plants

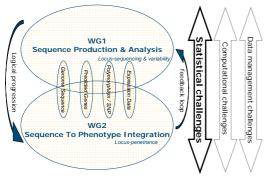
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Objective of EU COST Action TD0801

Establish a **network** of European researchers developing and/or using **efficient statistical** and **bioinformatics tools** and **strategies** to produce, assemble, analyze, and integrate high-throughput **Next Generation Sequence** data to increase our **knowledge** of crop and model **plant system**.

Figure 1: Two Working Groups of the COST Action and their putative interactions



Box 1: Next-Generation Sequencing (NGS) Technology platforms

Embodying a complex engineering of enzymology, chemistry, high-resolution optics, hardware and software. Providing significant time savings and minimal requirement for associated equipment a streamlined sample preparation prior to DNA sequencing. Examples are Roche/454 FLX, Illumina Genome Analyzer; and ABI SOLiD System.

This NGS technology will change the way on

- (Re-) Sequencing of (fragments of) genomes
- Profiling transcriptomes
- Elucidating DNA protein interactions
- Studying Epigenomics, Metagenomics, etc.

This EU COST Action Action is open to global participation – ready to collaborate with other ongoing initiatives outside Europe. The end date of the Action is anticipated to be early 2013. After installation of a Management Committee, Working Groups will be enforced that will organize the activities within the Action. *Facts and figures:* www.cost.esf.org/domains actions/fa/Actions/statistical challenges

Box 2: Uniqueness of plants

- repeat density in plants
- complications of polyploidy
- Immortalized (inbred/mutant) populations
- Extensive phenotyping in wide range of conditions

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