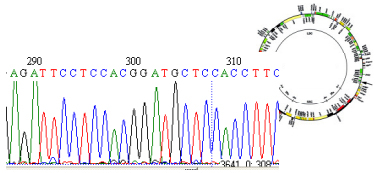
Complete cpDNA genome sequencing
→ 7 accessions + 1 out-group



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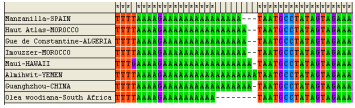
Polymorphism detection:
SNPs and length variants



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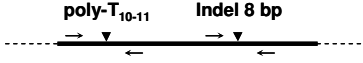
Marker development
(primers design for 64 loci)



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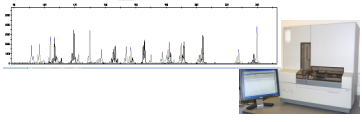
Screening of polymorphic loci on a set
of Mediterranean olive accessions



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Large scale genotyping (e.g. multiplex
PCR for microsatellites and indels)