Number of individuals in each map, project the variants belong to and methods used for CNV detection.

| Мар | Abyzov | Handsaker | Zarrei | Sudmant (Science) | Sudmant (Nature) |
|-------------|-------------------------------------|-------------------------------------|---------------------------------|----------------------|----------------------------------|
| Individuals | 1092 | 849 | 2647 | 236 | 2504 (from 26 populations) |
| Project | 1000 Genomes Project, Phase 1 | 1000 Genomes Project, Phase 1 | Meta analysis of DGV collection | | 1000 Genomes Project, Phase 3 |
| Methods | WGS, Split read | WGS, Read -depth | Multiple genome-wide techniques | WGS, Read-depth | WGS, Multiple algorithms |