|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | Number of 0.01 eQTLs (Best association per gene) | SNP genotyped and passed discovery threshold in 1KG | SNP genotyped and did not pass discovery threshold in 1KG | SNP not found in 1KG |
| CEU (Array) HM3  | 473 (498\*) | 241 (51.0%) | 221 (46.7%) | 11 (2.3%) |
| YRI (Array) HM3  | 489 (499\*) | 219 (44.8%) | 264 (54.0%) | 6 (1.2%) |
| CEU (RNA-Seq) HM3 | 965 (1071\*) | 327 (33.9%) | 626 (64.9%) | 12 (1.2%) |