| section | name | Europe | North America | Other |
| --- | --- | --- | --- | --- |
| Variants | p. nomenclature in variant list | 1.00 | 1.00 | 1.00 |
| Treatment | Therapeutic/Predictive information (if available) per variant | 0.92 | 0.96 | 1.00 |
| Variants | Gene symbol in variant list | 0.97 | 1.00 | 1.00 |
| Variants | Reported list of genetic variants | 0.93 | 1.00 | 1.00 |
|  | Gene list tested by assay | 0.96 | 0.90 | 0.93 |
|  | Tissue Source | 0.96 | 0.93 | 1.00 |
| Functional | Functional mechanism per variant (i.e. loss of function, activating) | 0.92 | 0.83 | 0.83 |
| Functional | Somatic variants classified for pathogenicity/oncogenicity | 0.88 | 0.83 | 0.86 |
| Overview | Diagnosis | 0.97 | 0.88 | 0.94 |
| Overview | Methods/Disclaimer: Assay limitations | 0.86 | 1.00 | 0.94 |
| Treatment | Diagnostic information (if available) per variant | 0.92 | 0.86 | 0.92 |
| Variants | Gene transcript per variant | 0.83 | 0.96 | 0.87 |
| Variants | c. nomenclature in variant list | 0.85 | 0.97 | 1.00 |
| Functional | Germline variants classified by ACMG/AMP criteria | 0.63 | 0.63 | 0.71 |
| Functional | Other Guidelines Used | 0.75 | 0.44 | 0.33 |
| Functional | Somatic variants classified by AMP/ASCO/CAP tiers | 0.64 | 0.93 | 0.64 |
| Functional | Variant prevalence in patient's tumor type | 0.42 | 0.50 | 0.80 |
| Overview | Methods/Disclaimer: Assay performance characteristics | 0.75 | 0.93 | 0.86 |
| Overview | Other disclaimers | 0.59 | 0.93 | 0.88 |
| Overview | Qualitative NGS Result Summary (i.e. This result is positive/negative) | 0.77 | 0.76 | 1.00 |
| Overview | Tumor purity/percentage | 0.96 | 0.70 | 0.87 |
| Treatment | Clinical Trial Information | 0.70 | 0.57 | 0.60 |
| Treatment | Databases/knowledgebases used to interpret variants | 0.82 | 0.79 | 0.85 |
| Treatment | Prognostic information (if available) per variant | 0.78 | 0.89 | 0.86 |
| Treatment | References used to interpret variant(s) | 0.86 | 0.97 | 0.77 |
| Treatment | Versions of databases used to interpret variants | 0.58 | 0.57 | 0.58 |
| Variants | Variant allele frequency per variant | 1.00 | 0.94 | 0.71 |
| Variants | g. nomenclature in variant list | 0.32 | 0.48 | 0.43 |
|  | List genes/regions with low coverage | 0.52 | 0.78 | 0.85 |