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Park K-J, et al.
A population study of inherited metabolic diseases

Supplemental Data Table S2. Variant calling criteria

Criteria	SNP	Indel
Minimum allele frequency	0.15	0.15
Minimum quality	10	10
Minimum coverage	20	20
Minimum coverage on either strand	3	3
Maximum strand bias	0.95	0.85
Minimum relative read quality	8.5	NA
Maximum common signal shift	0.25	NA
Maximum reference/variant signal shift (insertion)	NA	0.2
Maximum reference/variant signal shift (deletion)	NA	0.2

Abbreviation: NA, not applicable.