The Human Genetics Society of Australasia (HGSA) has reviewed the evidence and consulted with its expert members to develop the following recommendations to support best patient care and reduce the use of unnecessary or ineffective practices within a given clinical context.

1. Don’t use brain magnetic resonance imagery (MRI) for routine surveillance of asymptomatic neurofibromatosis type 1

2. Don’t undertake sequential testing for heterogeneous genetic disorders when targeted next generation sequencing (NGS) is available

3. Don’t undertake genetic testing for methylenetetrahydrofolate reductase (MTHFR), apolipoprotein E (APOE) and other such tests where the clinical utility for diagnostic purposes is extremely low

4. Don’t undertake carrier state testing for rare recessive disorders where a partner has a family history, the couple is non-consanguineous and there are no common causative mutations

5. Don’t undertake genetic testing when clinical diagnostic criteria exist and there are no reproductive or predictive testing implications

EVOLVE is a physician-led initiative to ensure the highest quality patient care through the identification and reduction of low-value practices and interventions. EVOLVE is patient-centred and evidence-based, with rigorous and transparent processes. Its focus is to stimulate clinical conversations – between colleagues, across specialties, and with patients – to ensure the care that’s delivered is the best for each patient.

EVOLVE is part of a worldwide movement to analyse medical practices and reduce unnecessary interventions. It is an initiative in partnership between the RACP and the Specialty Societies, Divisions, Faculties and Chapters.
1 Don’t use brain magnetic resonance imagery (MRI) for routine surveillance of asymptomatic neurofibromatosis type 1

Neurofibromatosis type I (NF-1) is a tumour disorder caused by the mutation of a gene on chromosome 17 that is responsible for control of cell division. It causes tumours along the nervous system that can grow anywhere in the body. Routine screening investigations are not recommended for the detection of the majority of complications associated with the condition. Baseline brain and spine MRI, and routine imaging of the chest and abdomen to identify asymptomatic tumours, do not influence management and should not be undertaken.

2 Do not undertake sequential testing for heterogeneous genetic disorders when targeted next generation sequencing (NGS) is available

A heterogeneous genetic disorder is one where the same disease or condition can be caused, or contributed to, by a number of different genes. The traditional strategy for genetic testing involves sequential sequencing of individual genes, selected according to the patient’s clinical presentation and family history. By contrast, next generation sequencing (NGS) involves the sequencing of millions of small fragments of DNA at the same time. Reductions in the cost of NGS now make it a more attractive solution for clinical diagnostic testing to identify the disease-causing mutation(s) in patients with genetically heterogeneous disorders than traditional sequential testing. In particular, the targeted NGS approach which restricts analysis to genes known to be implicated in a particular phenotype has been also successfully applied to heterogeneous disorders such as inherited peripheral neuropathy (IPN).

3 Do not undertake genetic testing for methylenetetrahydrofolate reductase (MTHFR), apolipoprotein E (APOE) and other such tests where the clinical utility for diagnostic purposes is extremely low

While genetic testing can help indicate susceptibility to particular genetic conditions, there are some conditions where the presence of particular alleles is neither necessary nor sufficient to cause the condition or where the alleles have a higher prevalence in the general population than the condition itself. This is the case for instance with apolipoprotein E as a genetic marker for Alzheimer’s disease and methylenetetrahydrofolate as a marker for venous thromboembolism.
4 Do not undertake carrier state testing for rare recessive disorders where a partner has a family history, the couple is non-consanguineous and there are no common causative mutations

With a rare recessive disorder, although the individual with the family history will have an increased risk of being a carrier, their unrelated partner will have a low general population risk. Therefore, their a priori combined risk of having a child with this rare recessive condition will generally be less than 1%. If the gene has no known common disease causative mutations then testing the unrelated partner for carrier status has low sensitivity and specificity.

5 Do not undertake genetic testing when clinical diagnostic criteria exist and there are no reproductive or predictive testing implications

Like other screening or diagnostic tests, genetic tests do not have inherent utility. It is the adoption of therapeutic or preventive interventions that influences health outcomes. If clinical diagnostic criteria already exist for the condition in question and there are no reproductive or other predictive testing implications as a result of definitively identifying a genetic cause for the condition, then this renders genetic testing unnecessary.

How this list was developed….  
A preliminary list was developed by the Lead Fellow which was then distributed to all the clinical geneticists in Australia who are all members of the Australasian Association of Clinical Geneticists (AACG), a special interest group of the HGSA. Following feedback the topic was revisited at a meeting of this group during the annual scientific conference of the HGSA, after which the list was finalised.
Human Genetics Society of Australasia – Top 5 recommendations

EVIDENCE SUPPORTING RECOMMENDATION 1

EVIDENCE SUPPORTING RECOMMENDATION 2
Antoniadi T, Buxton C, Dennis G. Application of targeted multi-gene panel testing for the diagnosis of inherited peripheral neuropathy provides a high diagnostic yield with unexpected phenotype-genotype variability. *BMC Medical Genetics* 2015; 16: 84
Ellard S, Lindsay H, Camm N. Practice guidelines for Targeted Next Generation Sequencing Analysis and Interpretation. *Association for Clinical Genetic Science Guidelines*, May 2014

EVIDENCE SUPPORTING RECOMMENDATION 3

EVIDENCE SUPPORTING RECOMMENDATION 4

EVIDENCE SUPPORTING RECOMMENDATION 5

DISCLAIMER: All reasonable care has been taken during the process of developing these recommendations. The health information content provided in this documents has been developed by the members of the Human Genetics Society of Australasia. The health information presented is based on current medical knowledge and practice as at the date of publication.