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Clinical scenario

What makes **PHARMACOGENETICS PSYCHIARTY** intriguing?

A significant percentage of patients do not benefit from antidepressant treatment and those who do, may incur severe side-effects.¹

The implicated long-term disability impacts on compliance, hospitalisations and healthcare costs. A 50-fold inter-individual variation in plasma levels is reported for drugs like amitriptyline² and since polypharmacy is common in psychiatry, interactions represent an additional concern.

Tricyclic antidepressants (TCAs) are often overlooked on claims of poor tolerability, although cheaper with a potential efficacy advantage. Much of the data available on TCAs derives from an era when genotyping studies were mostly unheard of. Research findings point towards a potential link between outcomes of amitriptyline therapy and the highly polymorphic CYP2D6 and CYP2C19.³ These two genes regulate the metabolism of approximately 25% of all prescription drugs⁴ implying that the result of one pharmacogenetic test is likely to be useful for a number of treatments.

Regulatory context

The FDA-approved drug label for amitriptyline states that CYP2D6 poor metabolizers may have higher plasma concentrations of TCAs, and suggests monitoring of plasma levels if this drug is co-administrered with a CYP2D6 inhibitor.⁵ There is the duty to guide in **interpretation** of this data so as to translate into actual **patient benefit**. It is understood that at the **EU** level, simple addition of information into the label is considered impractical.

For most medicinal products, genetic polymorphism consequences were noted after registration. To improve efficacy, avoid serious side-effects and promote cost effectiveness, genomically guided research should ideally not neglect currently marketed drugs but rather be undertaken to update the SmPC during the life cycle of the products. Prospective studies may not always be feasible due to older drugs having been replaced as first-line therapy, the need for large sample populations, and the lacking business incentive. Confirmatory evidence from case control studies, observational or epidemiological studies may serve as independent verification.⁶

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Research in this field may yield significant benefits for both the patient and the economy by helping change our focus from merely trying to find the best treatment for the typical patient. The 'ordinary' patient may, indeed, be uncommon.

Practical implications of pharmacogenetics in antidepressant treatment

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Is genotyping apt to determine the metaboliser status of patients receiving treatment for depression and translate biomarkers into personalised medicine? Can pharmacogenetics support the re-evaluation of tricyclic antidepressants with respect to effectiveness and safety, compared to newer, more expensive drugs 2 Should there be harmonisation at International, European and National level, in terms of what pharmacogenomic information is listed in drug labels?

OBJECTIVES

Investigate the incidence and impact of genetic polymorphisms in patients receiving treatment for depression in Malta -Evaluate genotype-phenotype association in relation to metaboliser status, measured blood drug/metabolite concentrations and clinical outcome -Analyse effect of regulatory initiatives in supporting the implementation of pharmacogenomics in drug surveillance and clinical practice -Pharmacoeconomic analysis of routine genotyping and worth of engineering point-of-care tests

Determine allele frequencies of both CYP2D6 and CYP2C19 and examine potential additive effects which would support combined dosing recommendations -Assess feasibility and clinician perception of pharmacogenetic-guided individualisation of treatment for depression, compared to the current standard of psychiatric care -

Analytical perspective

Genotyping data is often used to deduce enzyme hydroxylation capacity which nay at times be misinterpreted as the **metabolic phenotype**.⁷ The **ratio** between parent drug and relevant metabolites may allow better understanding of potentia correlations between genetics, enzyme functionality and clinical outcomes. In the case of TCAs, it is relevant to note, for example, that tricyclic hydroxymetabolites, like hydroxy-amitriptyline, are often excluded from therapeutic drug nonitoring even though these are associated with cardiotoxicity.⁸

Different DNA extraction kits allow comparison of blood vs saliva as source of DNA while distinct drug metabolism assays present the strengths and weaknesses of laboratory-based vs point-of-care technologies.⁹ CYP2D6 is a good example where comprehensive analytics are essential to avoid flawed conclusions in view of ie large number of polymorphisms to be analysed and the occurrence of copy mber variations.

References

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The case of amitriptyline

Metabolism, a key player in the dose-exposure-response chain, is a focus of promising genetically-oriented investigations underpinning the usefulness of genetic testing in the pharmacokinetic field. Several pharmaceutical companies are today integrating pharmacogenetic aspects in product labels and actionable genotype-guided dosing recommendations are available. **Conversely, many questions related to the implementation of personalised psychiatric therapy remain unanswered.**

STUDY DESIGN

Psychiatric outpatients receiving amitriptyline for depression • Free and informed consent • Blood, saliva and/or buccal cell samples • Depression and side-effects rating scales • Extensive genotyping and blood drug/metabolite monitoring • Results interpreted in line with gene/drug clinical practice guidelines Potential recommendations, if any, for treatment plan revision • Clinical outcomes observed on patient follow-up •

Retrospective Analysis

Past hospitalisations related to depression or adverse events from its treatment and the antidepressant switching strategies adopted. **Standard of Care**

Drugs with pharmacogenetic recommendations being prescribed at one time and the dose differences with respect to metaboliser status. Are non-extensive metabolizers being prescribed regular doses or are physicians empirically identifying patients with aberrant metabolism?

Prospective Correlations

The total range of variables encountered is included in the statistical analysis, combining both genetics and environmental factors such as age and age of onset, lifestyle, comorbidities, and comedications with special focus on concomitant CYP inhibitors, in attempt of explaining an appreciable percentage of variation.

Further Observations

Comparison of genotyping techniques, clinician disposition in implementing pharmacogenetic-guided recommendations, and the implications of the evolving regulatory context, particularly referral procedures assessed by the European Medicines Agency and review of pharmacogenomic data in national authorisations.

6. European Medicines Agency scientific guidelines on pharmacogenomics



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