



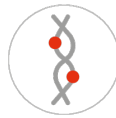
• Patient's name:	Dan Lawrence
• Received in laboratory	12-09-2024
• Genetic results date	12-11-2024

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# Hair Loss Treatment Test

## Results Report



## Genetic report

### **METHODOLOGY AND LIMITATIONS DISCLAIMER:**

Testing for genetic variation/mutation on listed genes was performed using Real-Time PCR with TaqMan® allele-specific probes on the QuantStudio 12K Flex. All genetic testing is performed by GX Sciences, LLC d/b/a Fagron Genomics US ("Fagron Genomics US") located at 807 Las Cimas Pkwy, Suite 145, Austin TX 78746. This test will not detect all the known alleles that result in altered or inactive tested genes. This test does not account for all individual variations in the individual tested. Test results do not rule out the possibility that this individual could be a carrier of other mutations/variations not detected by this gene mutation/variation panel. Rare mutations surrounding these alleles may also affect our detection of genetic variations. Thus, the interpretation is given as a probability. Therefore, this genetic information shall be interpreted in conjunction with other clinical findings and familial history for the administration of specific treatments. Patients should receive appropriate genetic counseling to explain the implications of these test results. The analytical and performance characteristics of this laboratory developed test (LDT) were determined by Fagron Genomics US's laboratory (Laboratory Director: James Jacobson, PhD) pursuant to Clinical Laboratory Improvement Amendments (CLIA) requirements (CLIA #: 45D2144988).

### **MEDICAL DISCLAIMER:**

This test was developed and its performance characteristics determined by Fagron Genomics US. It has not been cleared or approved by the FDA. The laboratory is regulated under CLIA and qualified to perform high-complexity testing. This test is used for clinical purposes. It should not be regarded as investigational or for research. The Reference SNP Cluster IDs (rsIDs) for the alleles being tested were obtained from the Single Nucleotide Polymorphism Database (dbSNP). These products are not approved by the Food and Drug Administration and are not intended to diagnose, treat, cure or prevent disease. These recommendations are for report purposes only and an individual is not required to use such products. These are recommendations only and do not replace the advisement of your own healthcare practitioner.

### **LEGAL DISCLAIMER:**

Report contents and report recommendations are created based on the consultation, advice, and direction of Dr. Gorana Kuka, the Medical Director for Fagron Genomics US TrichoTest protocols. Report contents and report recommendations are intended to be informational only. Report contents and report recommendations are not intended and should not be interpreted to make claims regarding the use, efficacy, or safety of products, formulas, and/or services listed herein. Only a doctor or other appropriately licensed health care practitioner can determine if a formula, product, or service described herein is appropriate for a specific patient. Sole risk for the use of all Fagron Genomics US lab test orders and test interpretation results rests with the reader. Implementation or experimentation with any supplements, herbs, dietary changes, medications, and/or lifestyle changes, etc. is done so at the patient's sole risk and responsibility and should be discussed with the patient or the patient's personal licensed healthcare practitioner prior to implementation. Fagron Genomics US and its affiliates, employees, associates, vendors, principals or partners, do not accept legal, moral, or ethical responsibility for any problems arising from experimentation with the information described in test results. Fagron Genomics US expressly reserves all legal rights and remedies in case of an inappropriate, negligent, or incorrect use or interpretation of the results of its tests.





# 1.

## Patient identification data



Ordering physician —●— **Riley Greene**  
Patient's name —●— **Dan Lawrence**  
Gender —●— **Male**  
Date of birth —●— **05-11-1974**  
Sample type —●— **Buccal Swab**  
Sample code —●— **TRI65317AA**  
Collection date —●— **12-02-2024**  
Received date —●— **12-09-2024**  
Genetic results date —●— **12-11-2024**



2.

Recommendation of the most suitable drugs and supplements

The **genetic test** uses an automated qualitative pharmacogenetic algorithm that analyzes the patient's genetic data and combines this information with relevant patient history to recommend the most suitable active ingredients. Next, we show on a color scale which compounds the algorithm recommends the most. The transition from white to dark green indicates drugs from least recommended to most recommended. Medications blocked due to intolerances or contraindications are shown in red.

Anti-alopecic drugs

Prostaglandins	
• Minoxidil	53%
• Cetirizine Hcl	50%
• Latanoprost	44%

Antiandrogenic	
• Dutasteride	100%
• Finasteride	99%
• Spironolactone	100%
• Melatonin	75%

Anti-inflammatory	
• Clobetasol propionate	
• Triamcinolone acetonide	
• Hydrocortisone	
• Betamethasone dipropionate	
• Desonide	
• Fluocinolone acetonide	

Immunomodulator	
• Tacrolimus	

Hair care supplements

Circulation	
• Arginine	57%
• Caffeine	25%

Collagen synthesis	
• Cystine	



Vitamin, mineral and antioxidant supplements

Vitamin deficiency		Keratolytic	
• Vitamin B12 (Cyanocobalamin)	100%	• Tretinoin	50%
• Vitamin B7 (Biotin)	67%		
• Vitamin D	67%		
• Vitamin E (Tocoferol)	67%		
• Vitamin C (Ascorbic Acid)			
• Lysine			
• Vitamin B9 (Folate)			

Minerals	
• Zinc gluconate	100%
• Zinc acetate	100%
• Iron sulfate	67%
• Magnesium Gluconate	







3.

Complete data

Data from the medical questionnaire

Patient demographics

Gender

Male

Age (years)

50

Family history of alopecia

Siblings

Hair loss data


Type of alopecia

Androgenic alopecia


Grade of alopecia

Grade Vertex


Norwood-Hamilton Scale




Type I




Type II




Type III




Vertex




Type IV



Type V



Type VI



Type VII

Clinical examination

Amount of hair loss

Little bit

Complaints associated with alopecia

No

Patchy alopecia

No

10 / 24

3.

Complete data

Pharmacogenetic results

1. Anti-alopecic drugs

Treatment efficacy with prostaglandin inhibitors

Prostaglandin D2			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
GPR44-1	rs545659	CT	Genetic result: Predisposition to slightly higher GPR44 mRNA stability. Interpretation: Higher expression of the Prostaglandin D2 receptor 2 (GPR44) receptor may lead to increased responsiveness to prostaglandin D2. Treatment recommendation: Consider treatment with Prostaglandin D2 inhibitors (e.g., Cetirizine, Prostaquinon) at standard doses.
GPR44-2	rs533116	CT	Genetic result: Predisposition to slightly higher GPR44 mRNA stability. Interpretation: Higher expression of the Prostaglandin D2 receptor 2 (GPR44) receptor may lead to increased responsiveness to prostaglandin D2. Treatment recommendation: Consider treatment with Prostaglandin D2 inhibitors (e.g., Cetirizine, Prostaquinon) at standard doses.

Latanoprost			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
PTGFR-1	rs6686438	GG	Genetic Result: This prostaglandin F receptor (PTGFR) variant may be associated with significantly lower probability of treatment efficacy with latanoprost (prostaglandin analog). Interpretation: Prostaglandin F receptor (PTGFR) variants are related with Latanoprost treatment efficacy (prostaglandin analog) . Treatment recommendation: Consider the use of concomitant or alternative therapies.
PTGFR-2	rs1328441	CT	Genetic Result: This prostaglandin F receptor (PTGFR) variant may be associated with an intermediate probability of treatment efficacy with latanoprost (prostaglandin analog). Interpretation: Prostaglandin F receptor (PTGFR) variants are related with Latanoprost treatment efficacy (prostaglandin analog) . Treatment recommendation: Consider treatment with Latanoprost at standard or higher doses.
PTGFR-3	rs10782665	GT	Genetic Result: This prostaglandin F receptor (PTGFR) variant may be associated with an intermediate probability of treatment efficacy with latanoprost (prostaglandin analog). Interpretation: Prostaglandin F receptor (PTGFR) variants are related with Latanoprost treatment efficacy (prostaglandin analog) . Treatment recommendation: Consider treatment with Latanoprost at standard or higher doses.

Treatment efficacy with minoxidil

Minoxidil			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
PTGES2	rs13283456	CT	Genetic Result: Predisposition toward slightly reduced PGE2 levels. Interpretation: Lower PTGES2 activity may result in significantly lower PGE2 levels which is related to alopecia development. Minoxidil promotes hair growth by increasing the production of PGE2 via PTGES2. Treatment recommendation: Consider oral, topical, or combination minoxidil treatment at standard doses to stimulate PGE2 production.
SULT1A1	rs1042028	AG	Genetic result: Predisposition to slightly reduced SULT1A1 activity and therefore less conversion of minoxidil into its active metabolite minoxidil sulfate. Interpretation: Minoxidil Sulfotransferase Enzyme (SULT1A1) variants predict response to minoxidil treatment. Treatment recommendation(s): Consider increasing the dosage or frequency of administration of oral or topical minoxidil. Consider combining oral and topical minoxidil treatments concurrently. Consider the use of a SULTA1A enzyme activity enhancing agent (e.g. tretinoin) compounded with topical minoxidil.

Treatment efficacy with glucocorticoid anti-inflammatories

Glucocorticoides			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
GR-alpha	rs6198	CT	Genetic result: Predisposition to moderate resistance to glucocorticoid anti-inflammatory treatments. Interpretation: Glucocorticoid Receptor (GR or NR3C1) variants are associated with resistance or sensitivity to corticosteroids. Treatment recommendation: If glucocorticoid anti-inflammatory treatment is used, doses should be slightly increased or an alternative treatment with non-glucocorticoid anti-inflammatory drugs should be chosen.

Treatment efficacy with antiandrogenics

17-α estradiol			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
CYP19A1	rs2470152	AA	Genetic result: Predisposition to significantly reduced CYP19A1 expression. Interpretation: Lower Aromatase (CYP19A1) activity may be associated with reduced conversion of testosterone into estrogens and higher conversion into DHT (a known hair growth inhibitor). Treatment recommendation: Consider treatment with 17-α Estradiol (an aromatase inducer) at higher doses. Consider adjuvant anti-androgen therapies and additional modalities.

Dutasteride			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
SRD5A1	rs39848	CC	Genetic result: Predisposition significantly increased SRD5A1 activity. Interpretation: Steroid 5α-Reductase 1 (SRD5A1) variants are associated with increased SRD5A1 activity leading to increased DHT levels and hair growth inhibition. Treatment/dosage: Treatment with Dutasteride at higher doses is recommended.

Finasteride			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
SRD5A2	rs523349	CG	Genetic result: Predisposition to slightly increased SRD5A2 activity. Interpretation: Steroid 5α-Reductase 2 (SRD5A2) variants are associated with increased SRD5A2 activity leading to increased DHT levels and hair growth inhibition. Treatment recommendation: Consider treatment with traditional doses of oral finasteride. Consider topical finasteride. Consider adding additional anti-androgen therapies or other modalities (e.g., pharmaceutical, phototherapy, nutraceutical, regenerative, etc.).

2. Hair care supplements

Vasodilatation and blood circulation

Circulation stimulators			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
ACE	rs4343	AG	Genetic result: Predisposition to slightly increased ACE activity. Interpretation: Increased Angiotensin-converting enzyme (ACE) activity may be associated with increased plasma levels of Angiotensin II, an extremely potent vasoconstrictor. Treatment recommendation: Consider normal doses of circulation stimulators such as Minoxidil, caffeine, Ginkgo Biloba, Ginseng or Arginine.

Collagen synthesis

Hair strengthening supplements			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
COL1A1	rs1800012	CC	Genetic result: Predisposition to normal COL1A1 expression. Interpretation: Increased COL1A1 expression is correlated with androgenetic alopecia and may be associated with a higher degree of collagen molecule instability due to an altered ratio of collagen α1/α2. Treatment recommendation: While hair strengthening composites may be of benefit, the SNP analysis does not indicate a relative indication nor contraindication for their use.

Reduction of IGF-1 levels

Hair strengthening supplements			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
IGF1R	rs2229765	AA	Genetic result: Predisposition to reduced IGF-1 levels. Interpretation: Insulin-like growth factor-I (IGF-I) variants are associated with lower plasma IGF-1 levels leading to hair loss. Treatment recommendation: A treatment with Igrantine-F1 and TrichoXidil (IGF-1 inducers) at high doses would be recommended.

3. Vitamin, mineral and antioxidant supplements

Vitamins

Vitamin A			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
CRABP2	rs12724719	AG	Genetic Result: This CRABP2 variant is associated with slightly lower retinoic acid (vitamin A) intracellular transport. Interpretation: Cellular retinoic acid-binding protein 2 (CRABP2) variants are associated with lower retinoic acid (vitamin A) intracellular transport. Treatment recommendation: Consider nutritional supplementation with normal doses of vitamin A.
Vitamin B7			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
BTD	rs13078881	CG	Genetic Result: This BTD variant suggests a predisposition to slightly reduced Biotidinase activity. Interpretation: Reduced biotinidase activity may result in low biotin levels due to significantly lower biotin (vitamin B7) uptake from the gut. Treatment recommendation: Consider supplementation with normal doses of B vitamins.
Vitamin C			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
SLC23A1	rs33972313	CC	Genetic result: Predisposition to higher vitamin C serum level. Interpretation: Solute carrier family 23 member 1 (SLC23A1) variants are associated with lower serum concentration of vitamin C. Treatment/dosage: SNP analysis does not indicate the necessity to supplement with vitamin C. Test for serum levels of vitamin C.
Vitamin B9			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
MTHFR	rs1801133	GG	Genetic result: No predisposition to folate deficiency. Interpretation: Methylene tetrahydrofolate reductase (MTHFR) variants are associated with risk of folate deficiency. Treatment/dosage: SNP analysis does not indicate the necessity to supplement with folate. Test serum levels of folate prior to supplementation.

Vitamin D			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
GC	rs2282679	GT	Genetic result: Predisposition to slightly lower vitamin D serum level. Interpretation: Vitamin D-binding protein (GC or DBP) variants are associated with lower vitamin D serum level. Treatment/dosage: Supplementation should be considered. Test serum levels of vitamin D prior to supplementation.

Vitamin B12			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
FUT2	rs602662	AG	Genetic result: Predisposition to low vitamin B12 serum level. Interpretation: Galactoside 2-alpha-L-fucosyltransferase 2 (FUT2) variants are associated lower vitamin B12 serum level. Treatment/dosage: Supplementation with vitamin B12 is highly recommended.

Vitamin E			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
ZPR1	rs964184	CG	Genetic result: Predisposition to slightly lower serum tocopherol levels. Interpretation: Zinc Finger Protein ZPR1 variants are associated with low serum alpha-tocopherol (vitamin E) levels. Treatment/dosage: Vitamin E supplementation should be considered. Test serum levels of vitamin E prior to supplementation.

Antioxidants

Antioxidants			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
NQO1	rs1800566	GG	Genetic result: Predisposition to normal NQO1 enzyme activity. Interpretation: NAD(P)H dehydrogenase [quinone] 1 (NQO1) variants are associated with lower NQO1 enzyme activity and may have less effective protection against oxidative stress. Treatment/dosage: SNP analysis does not indicate the necessity to supplement with antioxidants. Test serum levels of selenium prior to supplementation.

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## Minerals

Magnesium			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
MUC1	rs4072037	TT	Genetic result: Predisposition to higher magnesium serum level. Interpretation: Mucin 1, cell surface associated (MUC1) variants are associated with lower magnesium serum level. Treatment/dosage: SNP analysis does not indicate the necessity to supplement with magnesium. Test serum levels of magnesium prior to supplementation.

Zinc sulfate			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
SLC30A3	rs11126936	GG	Genetic result: Predisposition to lower serum zinc level. Interpretation: Solute carrier family 30 member 3 (SLC30A3) variants are associated with lower serum zinc level. Treatment/dosage: Supplementation with Zinc sulfate would be highly recommended. Test serum levels of zinc prior to supplementation.

Iron			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
TMPRSS6	rs855791	AG	Genetic result: Predisposition to slightly reduced serum levels of transferrin and iron. Interpretation: Transmembrane protease, serine 6 (TMPRSS6 or matriptase-2) variants are associated with decreased serum levels of transferrin and iron. Treatment/dosage: Supplementation should be considered. Test serum levels of iron prior to supplementation.

Selenium			
Gene	SNP (transition)	Patient genotype	Pharmacogenetic result
DMGDH	rs921943	CC	Genetic result: Predisposition to lower selenium serum level. Interpretation: Dimethylglycine dehydrogenase (DMGDH) variants are associated with low selenium serum level. Treatment/dosage: Selenium supplementation should be considered. Test serum levels of selenium prior to supplementation.





## 4. Methodology

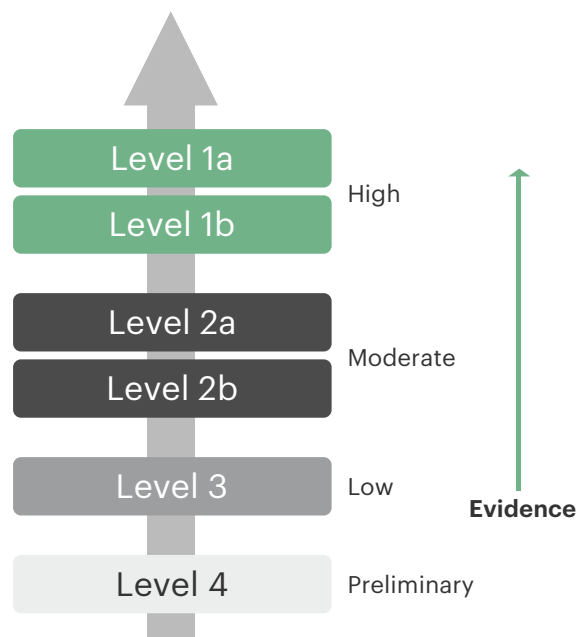
### How were the genetic variants studied selected and evaluated?

The **genetic test** was developed by a multidisciplinary team of medical doctors, pharmacists, geneticists, and programmers, following the highest quality standards. In particular, an expert team specialized in the curation of genetic variants reviewed each variant to ensure that selection, interpretation and impact of variants in the algorithms are based on the highest scientific evidence. Relevant patient's anamnesis (intolerances, diseases, medication, blood pressure, among others) that can affect recommendations was taken into account through medical questionnaires elaborated by health professionals.

- **Level 1A:** Annotation for a variant in medical society-endorsed or implemented in a major health system.
- **Level 1B:** Annotation for a variant where the preponderance of evidence shows an association. The association must be replicated in more than one cohort with significant p-values, and preferably will have a strong effect size.
- **Level 2A:** Annotation for a variant that qualifies for level 2B where the variant is within a Very Important known gene, so functional significance is more likely.
- **Level 2B:** Annotation for a variant with moderate evidence of an association. The association must be replicated but there may be some studies that do not show statistical significance, and/or the effect size may be small.
- **Level 3:** Annotation for a variant based on a single significant (not yet replicated) study or annotation for a variant evaluated in multiple studies but lacking clear evidence of an association.

- **Level 4:** Annotation based on a case report, non-significant study or in vitro, molecular or functional assay evidence only.

Only variants from level 1a to 2b were selected.



### How was this test performed?

DNA was extracted from the buccal swab sample provided and was analyzed by our clinical analysis laboratory. DNA was extracted using the KingFisher Flex® robotic extraction system (Thermo Fisher Scientific). The study of the genetic variants was carried out using a custom-designed microfluidic card to measure for the chemiluminescent detection of each of them using TaqMan® technology. TaqMan® technology for genotyping testing is proven and widely used in clinical and research settings. The sensitivity (detection limit) of this study is 99%.

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• Sample code: **TRI65317AA**

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## Genetic test algorithm

The **genetic test** qualitative pharmacogenetic algorithm analyzes single nucleotide polymorphisms (SNPs) associated with metabolic pathways involved in alopecia predisposition and treatment and combines this data with relevant patient history to predict treatment responses and recommends the most appropriate active ingredients.

The **genetic test** is an in vitro diagnostic medical device developed by **Fagron Genomics** and marketed under the CE-IVD mark in conformity with European Directive 98/79/EC and the transitional provisions (article 130) of European Regulation 2017/746.



**Fagron Genomics S.L.U.,**

SRN: ES-MF-000001092

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## What are the limits of this report?

Each genetic marker tested is just one factor that predicts the likelihood of a particular outcome. However, the lifestyle, diet, and environment to which the patient is exposed may impact the expected outcomes. These external factors cannot be taken into account in this report.

The information in this report is not used to diagnose genetic diseases or abnormalities, as it does not predict the risk and likelihood of certain genetic outcomes. It is also not intended to diagnose or cure any disease. The **genetic test** is intended to assist health professionals in making patient-specific care decisions regarding the treatment or prevention of androgenetic alopecia, areata alopecia, and telogen effluvium.

Our clinical laboratory has standard and effective procedures to protect against technical and operational problems. However, problems may occur in the shipment to the laboratory or in the handling of the sample, including, but not limited to, damage to the sample, mislabeling, and loss or delay in receiving the sample. In such cases, the medical laboratory may need to request a new sample.

As with all medical laboratory tests, there is a small chance that the laboratory may provide inaccurate information.

It is the responsibility of the professional who requests a test from us to guarantee the interested party appropriate genetic counseling in accordance with Law 14/2007, of July 3, on Biomedical Research.

**Fagron Genomics S.L.U.** declines all responsibility derived from the use and interpretation of the results of our tests by the requesting health professional.

**Fagron Genomics S.L.U.** does not access data identifying the patient from whom the sample comes, so it is also the responsibility of the requesting professional to comply with the applicable data protection regulations.



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• Patient name: **Dan Lawrence**

• Date of birth: **05-11-1974**

• Sample code: **TRI65317AA**

• Received in laboratory: **12-09-2024**

• Genetic results date: **12-11-2024**

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