

Does IRF6 Work Alone?

Is IRF6 a lone actor, or one link in a chain of genes, and how would we tell from the patterns in patients?

CARRY FORWARD

A missense variant matters when its side-chain change disrupts a specific protein interaction or fold.

10-YEAR TAKEAWAY

IRF6 works in a regulatory network, so similar phenotypes can arise when different connected genes fail.

Cells differentiate	Development sculpts	Cells move	Development can go wrong
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1. Observe the analogy before naming the biology



Illustration: A relay team passes one signal through several runners

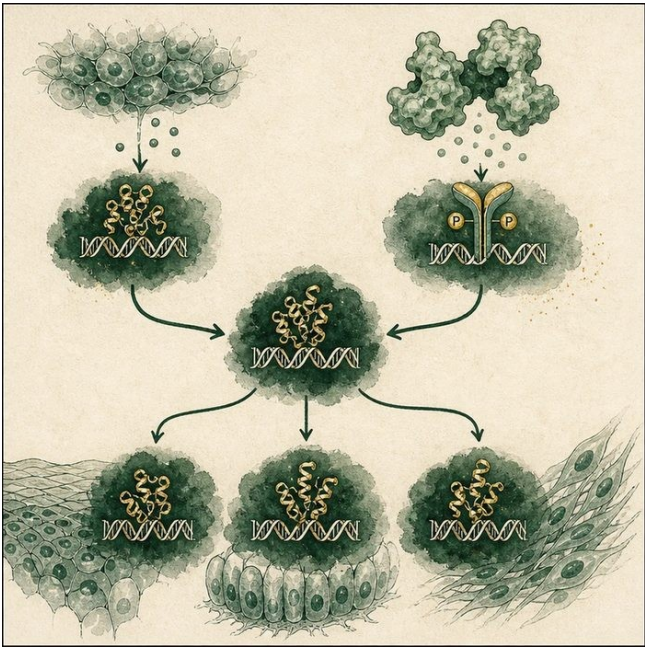
- 1. Who acts before the middle runner?
- 2. Which handoff could stop the same finish?
- 3. Why would two different failed runners create similar outcomes?

2. Turn observations into rules

- 1. Rule 1: Name upstream and downstream relationships. Explain it in your own words.
- 2. Rule 2: Different nodes can converge on one cell behavior. Explain it in your own words.
- 3. Rule 3: Network redundancy can soften or redirect a failure. Explain it in your own words.

Where the analogy stops: Gene networks branch, feed back, and vary by tissue, unlike a straight relay.

3. Map the rules onto the biomedical example



Biomedical teaching illustration: Place p63, IRF6, GRHL3, and KLF genes in a network

Analogy step	Biology step	How the relationship stays the same
Earlier runner	Upstream regulator	_____
Baton handoff	Transcriptional regulation	_____
Finish-line action	Epithelial differentiation and safe fusion	_____

GENETICS LESSON 13 | CASE FILE AND DECISION

4. Use the case evidence

ID	Finding supplied in the case	Why it matters
GEN13-E1	p63 and TGF-beta signaling influence IRF6 expression or activity in oral epithelium.	IRF6 has upstream regulators.
GEN13-E2	IRF6 promotes GRHL3, KLF4, and KLF17-related differentiation programs.	IRF6 connects to downstream targets.
GEN13-E3	GRHL3 variants can also cause Van der Woude-like phenotypes.	Different network nodes can converge on similar outcomes.

5. Make the clinical decision

You are explaining a negative IRF6 test in a family with lip pits and clefting. IRF6 sequencing is negative, but the phenotype remains strongly suggestive of the network.

- A. Consider other connected genes and broader testing.
- B. Conclude the phenotype is impossible.
- C. Invent an IRF6 variant.

Decision. Choose the next genetic strategy and cite network convergence.

Claim ceiling: You may broaden the candidate network. You may not diagnose a specific second gene without evidence.

Claim. State the choice you can defend.

Evidence. Use these labeled IDs: GEN13-E1 + GEN13-E2.

Reasoning. Explain why the evidence supports the claim and stays inside the claim ceiling.

6. Reduce the lesson to one lasting idea

Take-home. IRF6 works in a regulatory network, so similar phenotypes can arise when different connected genes fail.

1. What do upstream and downstream mean?
2. Why can different genes produce similar phenotypes?

Glossary

Term	Plain-English definition
gene regulatory network	The web of genes that turn each other on and off to control a process such as palate fusion.
transcription factor	A protein that binds DNA and turns specific genes on or off, controlling what a cell becomes and does.
upstream	Genes or events that come earlier in a pathway and control what happens later, downstream.
downstream	Describing genes or events that come later in a pathway, controlled by something acting earlier, or upstream, of them.
redundancy	When more than one gene can perform the same job, so losing one is partly covered by another and the effect is softened.

Next lesson unlock

We placed IRF6 in a network with p63 above it and GRHL3 and the KLFs alongside and below. But a network of genes is just instructions. What does this network actually make the cells of the embryo do so the lip and palate can physically close? We chase that next.

The journal links on the lesson page are optional. Everything required for this guided-notes set is already supplied here and in the case file.