

## Supplementary Material

### Identification of The Role of Wnt/ $\beta$ -catenin Pathway through Integrated Analyses and *in vivo* Experiments in Vitiligo

**Table S1** The Clinical Information of The Patients

Patient ID	Gender	Age	Duration	Progression	Classification	BSA <sup>a</sup>	Biopsy Area	Associated Disorders <sup>b</sup> or Family History
1	F	62Y	2Y	Stable	NSV	1%	Vulva	N
2	F	37Y	1Y	Stable	NSV	4%	Scalp	N
3	M	18Y	3Y	Stable	NSV	1.5%	Face	N
4	M	26Y	2Y	Stable	NSV	3%	Prepuce	N
5	F	22Y	5 Mon	Unstable	NSV	2%	Upper Limb	N
6	F	25Y	13Mon	Unstable	NSV	4%	Perianal Region	N

**Notes:** **a.** The severity of the disease is scored by body surface area(BSA) of depigmentation.

**b.** Associated disorders include psoriasis, alopecia areata, lupus erythematosus, rheumatoid

arthritis, diabetes and so on. All patients had not received vitiligo-related treatment prior to

surgical biopsy. **Abbreviations:** F, female; M, male; Y, years; Mon, months; NSV,

nonsegmental vitiligo; BSA, body surface area; N, none;

**Table S2** The Clinical Information of The Healthy Volunteer

Healthy Volunteer ID	Gender	Age	Biopsy Area	Associated Disorders <sup>a</sup> or Family History of Vitiligo
1	F	29Y	Cervical Region	N
2	F	28Y	Back	N
3	M	36Y	Cervical Region	N
4	M	48Y	Face	N
5	F	29Y	Back	N
6	M	32Y	Upper Limb	N

**Notes: a.** Associated disorders include psoriasis, alopecia areata, lupus erythematosus, rheumatoid arthritis, diabetes and so on.

**Abbreviations:** F, female; M, male; Y, years; N, none.

**Table S3** The q-PCR Reaction Conditions

Stages	Reps	Temperature	Time
Stage1	1	95°C	5min
Stage2	40	95°C	10sec
		60°C	30sec
Stage3	1	95°C	15sec
		60°C	60sec
		95°C	15sec

Abbreviations: min: minute; sec: second.

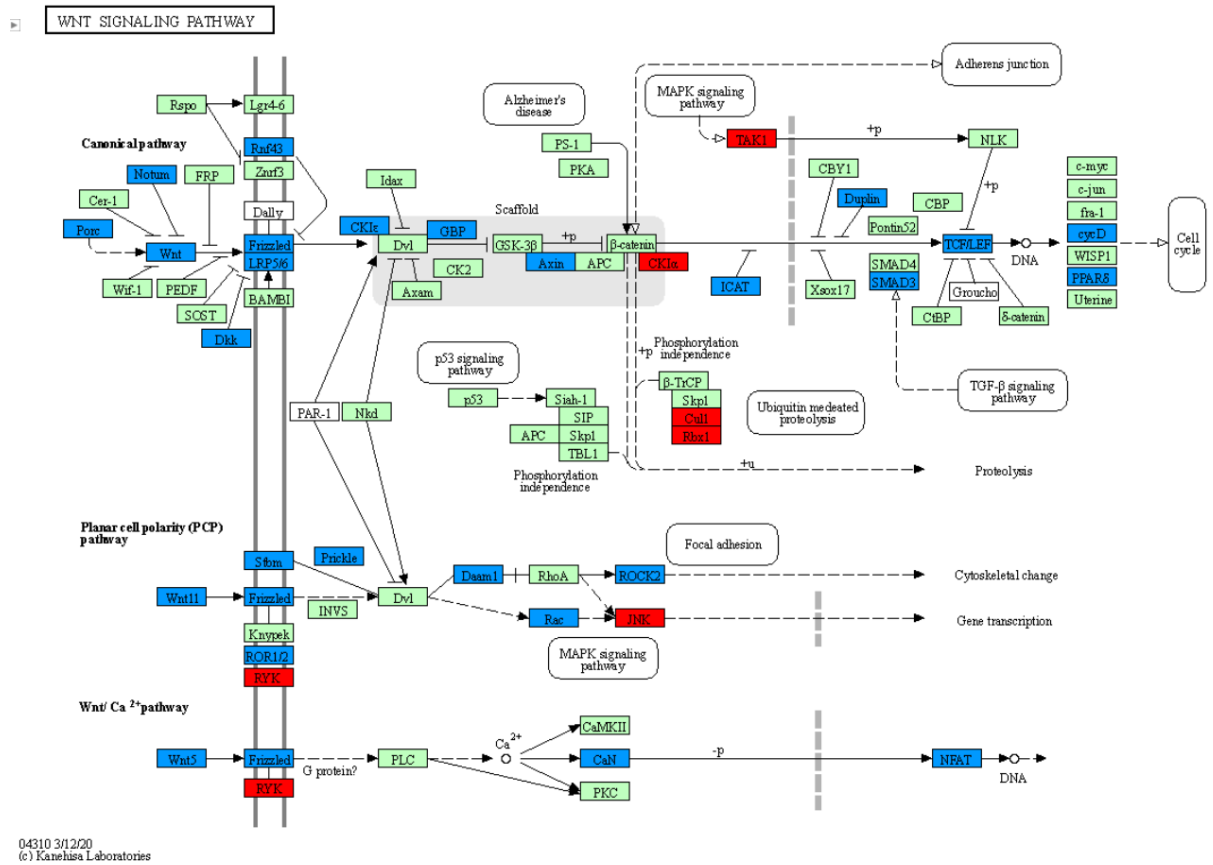
**Table S4** The Primer Information For q-PCR Analysis in Zebrafish.

Gene	Primer Sequence (5'to 3')
<i><math>\beta</math>-actin</i> <sup>1</sup>	Forward: GAAGGAGATCACCTCTCTTGCTC
	Reverse: GTTCTGTTTAGAAGCACTTCCTGTG
<i>tyr</i>	Forward: ACTACCGAGAGGCAGAGGTTTCATC
	Reverse: TTGGCGAACATTGGCGTGGAG
<i>mitf</i> <sup>2</sup>	Forward: TGTACAGCAATCATGCTCTTCC
	Reverse: GTCCCCAGCTCCTTAATTCTGTC
<i><math>\beta</math>-catenin</i> <sup>3</sup>	Forward: CAAGAGCAAGTAGCAGACATCG
	Reverse: CTGTGTGGAAGGTATCTGCATG
<i>lef1</i> <sup>4</sup>	Forward: CACGTTCTACACCTGTCCCTAA
	Reverse: AAGCAAGTGAGCGAGTGGACAT

**Source / reference of the primers:**

1. Wehner D, Cizelsky W, Vasudevaro MD, et al. Wnt/ $\beta$ -catenin signaling defines organizing centers that orchestrate growth and differentiation of the regenerating zebrafish caudal fin. *Cell Rep.* 2014;6(3):467-81. doi: 10.1016/j.celrep.2013.12.036.
2. Lister JA, Capper A, Zeng Z, et al. A conditional zebrafish MITF mutation reveals MITF levels are critical for melanoma promotion vs. regression *in vivo*. *J Invest Dermatol.* 2014;134(1):133-140. doi: 10.1038/jid.2013.293.

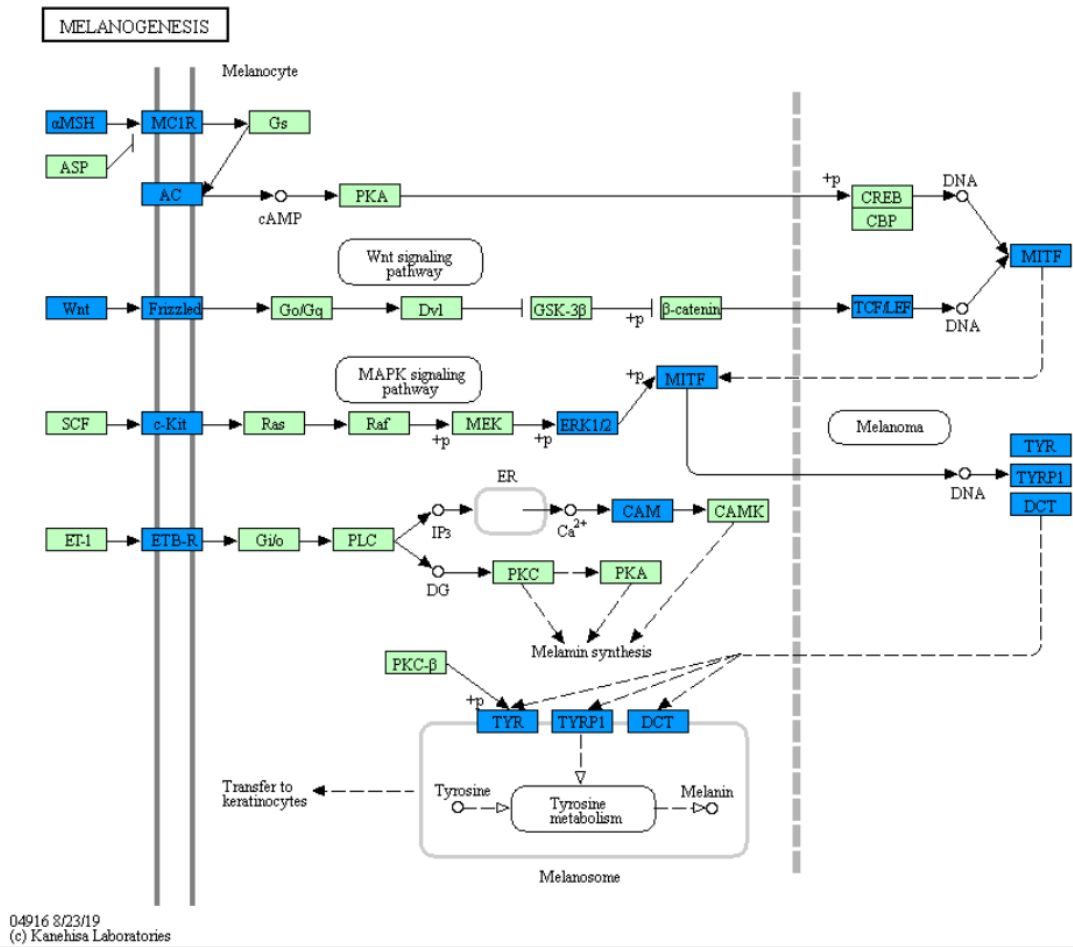
3. Gorsuch RA, Lahne M, Yarka CE, Petravick ME, Li J, Hyde DR. Sox2 regulates Müller glia reprogramming and proliferation in the regenerating zebrafish retina via Lin28 and *Ascl1a*. *Exp Eye Res.* 2017;161:174-192. doi: 10.1016/j.exer.2017.05.012.
4. Yin A, Korzh S, Winata CL, Korzh V, Gong Z. Wnt signaling is required for early development of zebrafish swimbladder. *PLoS One.* 2011;6(3):e18431. doi: 10.1371/journal.pone.0018431.



**Figure S1** DEGs enriched in Wnt signaling pathway. Red represented upregulated DEGs and

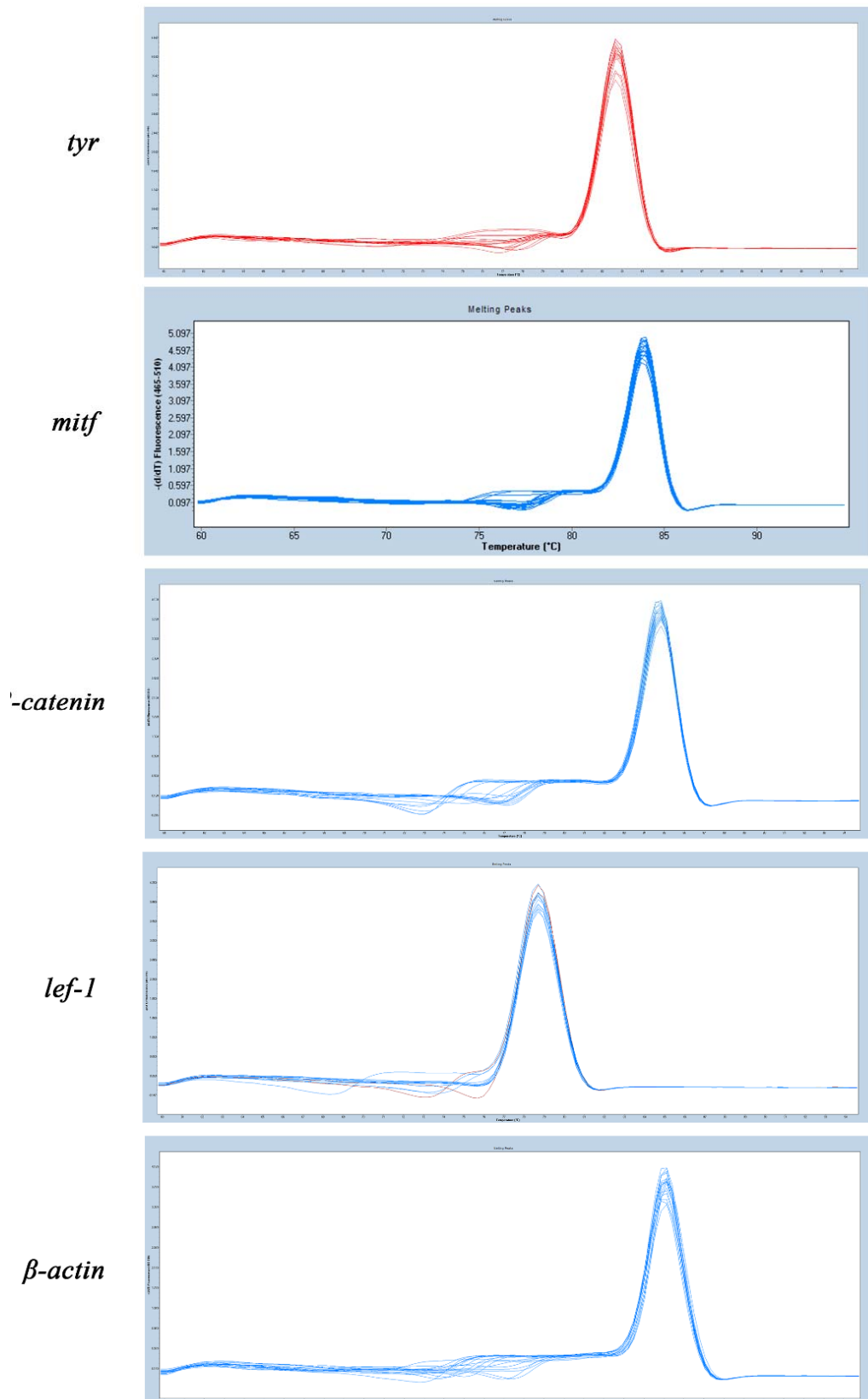
Blue represented downregulated DEGs. DEGs, differentially expressed genes.

Reproduced from: Kanehisa M, Furumichi M, Sato Y, Ishiguro-Watanabe M, Tanabe M. KEGG: integrating viruses and cellular organisms. *Nucleic Acids Res.* 2021;49(D1):D545-D551. doi: 10.1093/nar/gkaa970



**Figure S2** DEGs enriched in Melanogenesis signaling pathway. Red represented upregulated DEGs and Blue represented downregulated DEGs. DEGs, differentially expressed genes.

Reproduced from: Kanehisa M, Furumichi M, Sato Y, Ishiguro-Watanabe M, Tanabe M. KEGG: integrating viruses and cellular organisms. *Nucleic Acids Res.* 2021;49(D1):D545-D551. doi: 10.1093/nar/gkaa970



**Figure S3** Melt-curve analysis of *tyr*, *mitf*,  $\beta$ -*catenin*, *lef-1* and  $\beta$ -*actin* genes. The melting curves are single peaks, indicating that the products of q-PCR are specific.